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Sequence:

Sequence 44, 7 Sequence 338, Sequence 8, Ag

Sequence 6, Al Sequence 10, 3 Sequence 927,

Sequence 1 Sequence 1 Sequence 1

Sequence 1 Sequence 1 Sequence 1

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WS-09-300-958A-27

WS-09-300-958A-27

WS-09-300-958A-27

Sequence 27, Application US/09300958A

Patent No. 64531-100

Patent No. 64531-100

APPLICANT: McClelland, Michael

APPLICANT: McClelland, Michael

APPLICANT: McClelland, Michael

APPLICANT: McClelland, Michael

APPLICANT: Weduced Complexity Nucleic Acid Targets and Methods of

TITLE OF INVENTION: UBing Same

FILE REFERENCE: 1999-04-27

CURRENT APPLICATION NUMBER: G0/098,070

PRIOR PAPLICATION NUMBER: G0/108,624

PRIOR APPLICATION NUMBER: G0/118,624

PRIOR APPLICATION NUMBER: G0/118,624

NUMBER OF SEQ ID NOS: 85

SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                       Sequence 1, Appli
Sequence 3, Appli
Sequence 1, Appli
Sequence 12, Appli
Sequence 12, Appli
Sequence 13, Appli
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Mismatches:
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Matches:
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Best Local Similarity:
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Sequence 4, Appli
Sequence 1, Appli
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Sequence 2, App
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                        - nucleic search, using frame_plus_p2n model
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                                                                                                                                                                                                                                                                                                                                                                              398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    160 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyArgGlnAlaSerProTyrTyr 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         200 ThralaThrProGlnSerSerHisAlaSerAspSerGlyGlySerAspValAspLeuAsp 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         220 LeuThrGluSerLysValPheProArgAspAspPheThrAspTyrLysLysGlyGluPro 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              260 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheIleArg 279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        320 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGlu 339
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                                                                             GluAspProThrLeuAlaProAlaProPro---ThrThrPheGlyThrGluAspLeuVal 39
                                                                                                                                                             LeuThrieuAsnAsnGlnGlnMetThrieuGluGlyProGluLysAlaSerTrpThrSer 59
                                                                                                                                                                                                                                                                                                                                LysAsnLysTyrAspAlaSerSerlleAspPheSerArgCysAsnMetAspGlyAlaThr 99
                                                                                                          159 GAGGACTCCACCCTGGCTCTGTTCCCCCTGCTGCCACTTTGGGGCCGATGACTTGGTA
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AlaAlaThrCysGluIleSerAsnValPheSerAsnTyrPheAsnAlaMetTyrSerSer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 120 HisAlaGlnLeuArgAspLeuThrSerAsnSerSerAspGluLeuSerTrpIleIleGlu 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LeuLeuGluLysAspGlyMetSerPheGlnGluSerLeuGlyAspLeuGlyProPheAsp 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        519 CTGCTGGAGAAGGAATGGCATTGCAGGAGGCCCTA---GACCCAGGGCCCTTTGAC 575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         40 LeuThrieuAsnAsnGlnGlnMetThrieuGluGlyProGluLysAlaSerTrpThrSer 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | GluArgProGlnPheTrpSerLysThrGlnValLeuGluTrpIleSerTyrGlnValGlu 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 AlaAlaThrCysGluIleSerAsnValPheSerAsnTyrPheAsnAlaMetTyrSerSer
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ORGANISM: Homo sapiens
FRATURISM: Homo sapiens
NAME/KBY: CDS
LOCATION: (96)...(1211)
OTHER INFORMATION: Protein.
                                                                                                 Sequence 4, Application US/09570593
Patent No. 656663
GENERAL INFORMATION:
APPLICANT: Raifmann, Joerg
APPLICANT: Antiman, Joerg
APPLICANT: Harrow, Greg
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: 2000-05-12
FILE REPERBNCE: 2100-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT PILING DATE: 1200-1556
NUMBER OF SEQ ID NOS: 13
NUMBER OF SEQ ID NOS: 13
SOTWARE: FastSEQ for Windows Version 4.0
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Matches:
Conservative:
Mismatches:
Indels:
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Best Local Similarity:
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160 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyArgGlnAlaSerProTyrTyr 179
                            180 CysSerThrTyrGlyProGlyAlaProSerProGlySerSerAspValSerThrAlaArg 199
                                                                                                                             636 CCCGGCAGCTGTGCCCCAGGAGCCCCCTCGCCAGCTCTGACGTCTCCACCGCAGGG 695
                                                                                                                                                                                200 ThralaThrProGlnSerSerHisAlaSerAspSerGlyGlySerAspValAspLeuAsp 219
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Patent No. 5789200
GENERAL INFORMATION:
APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: 4 No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
CHTY: King of Prussia
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION: 514
APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM 486
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STATE: PA
COUNTRY:
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ATTORNEY/AGENT INFORMATION:

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Conservative:
Mismatches:
Indels:
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        REGISTRATION NUMBER: 34,344
REFERENCE/DOCKET NUMBER: ATG 50024
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270 4026
INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
LENGTH: 1920
TYPE: Nucleic Acid
STRANDEDNESS: Single
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1707.00
92.72%
86.79%
86.21%
William T. Han
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Best Local Similarity:
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| Sequence 282, Application US/09389681A |
| GENERAL INFORMATION: |
| APPLICANT: Vuqui, Jiang |
| APPLICANT: Mitcham, Jennifer L. |
| APPLICANT: Mitcham, Jennifer L. |
| APPLICANT: Xu, Jiangchun COMPOSITIONS FOR THE TREATMENT AND |
| TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE |
| TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE |
| CURRENT FILING DATE: 1999-09-02 |
| CURRENT FILING DATE: 1999-09-02 |
| NUMBER OF SEQ ID NOS: 463 |
| SOFTWARE: FastSEQ for Windows Version 3.0 |
| SERVID NO 282 |
| LENGTH: 502 |
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240 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys 259
                                                                              260 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg 279
                                                                                                              895 CTCGAGGGCAAGAAGAAGCACGCCCAGAGGCACCCACCTGTGGGAGTTCATCCGG 954
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                            835 AACCACGGGAAGCGGAAACGAGGCCCCGGAAAGCTGAGCAAAGAGTACTGGGACTGT
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Matches:
Conservative:
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93.33$
90.30$
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ORGANISM: Homo sapiens
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Best Local Similarity:
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US-09-389-681-282
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Pred. No.:
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186 CGGAAACGAGGCCGCGGAAAGCTGAGCAAAGAGTACTGGGACTGTCTCGAGGGCAAG 245
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Patent No. 6528054
GENERAL INFORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Miccham, Jennifer L.
APPLICANT: Miccham, Jennifer L.
APPLICANT: Miccham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: COMPOSITIONS
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILER REFERENCE: 210121.470C8
CURRENT APPLICATION NUMBER: US/09/620,405B
CURRENT APPLICATION NUMBER: US/09/620,405B
CURRENT APPLICATION NUMBER: US/09/620,405B
SOFTWARE: FREESE for Windows Version 3.0
SEQ ID NO 282
LENGTH: 502
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Matches:
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Query Match:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Vigiu, Jiang
APPLICANT: Vigiu, Jiang
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C2
CURRENT APPLICATION WUMBER: US/09/339,338A
CURRENT FILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 315
SOFTWARE: FastSEQ for Windows Version 3.0
264 LysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheIleArgAspIleLeuIle
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ORGANISM: Homo sapiens
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LENGTH: 502
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APPLICANT: Dillon, Davin C.
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun C.
APPLICANT: Xu, Jiangchun C.
APPLICANT: Xu, Jiangchun C.
APPLICANT: NYENTON: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210121.47004
CURRENT FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 4774
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 282
FENDERM: 602
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Sequence 282, Application US/09433826B

; Patent No. 6579973

; GENERAL INFORMATION:
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426 ACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAGATCCTGGAACGG 485
      324 ThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGluIleLeuGluArg 343
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Matches:
Conservative:
Mismatches:
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                                                                                                                                                                                                              APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels:
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APPLICATION NUMBER: US/09/009,913
FILING DATE: 21-JAN-1998
                                                                                                                                                                 ; Sequence 2, Application US/09009913
; Patent No. 6087485
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: SE
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
                                                                      344 ValAspGlyArgArg 348
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pair
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        , MOLECULE TYPE: CDNA
US-09-009-913-2
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                               USA
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                                                                                                                                 RESULT 9
US-09-009-913-2
                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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324 ThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrTysBrgGlulleLeuGluArg 343
                                                                      66 CGGAGCTCCCACTCCTCAGACTCCGGTGGAAGTGACGTGGACCTGGATCCCACTGATGGC 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       184 GlyProGlyAlaProSerProGlySerSerAspValSerThrAlaArgThrAlaThrPro 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    224 LysvalpheproArgAspAspPheThrAspTyrLysLysGlyGluProLysHisGlyLys 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     264 LysserLysHisAlaProArgGlyThrHisLeuTrpGluPheileArgAspIleLeuIle 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 284 HisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGluGlyValPheLys 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       204 GlnSerSerHisAlaSerAspSerGlyGlySerAspValAspLeuAspLeuThrGluSer 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    244 ArglysArgGlyArgProArglysLeuSerLysGluTyrTrpAspCysLeuGluGlyLys 263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  246 AAGAGCAAGCACGCCCAGAGGCACCCACCTGTGGGAGTTCATCCGGGACATCCTCATC 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  306 CACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAAGGCGTCTTCAAG 365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 304 PheleuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysAsnSerAsnMet 323

    6 GGCGCAGGAGCCCCCCCCCCGGCAGCTCTGACGTCTCCACCGCAGGGACTGCTTCT

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Matches:
Conservative:
Mismatches:
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SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 282
LENGTH: 502
                                                                                                               344 ValAspGlyArgArg 348
                                                                                                                                              486 GTGGATGGCCGCGA 500
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93.33%
90.30%
40.56%
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; ORGANISM: Homo sapiens
US-09-604-287A-282
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Best Local Similarity;
                                                                                                                                                                                           US-09-604-287A-282
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Pred. No.:
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141 -----AATGAGATCATGATTCTGGAAGGAGGTGTAATGAATCTCAACCCCGGCAAC 194
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195 AACCTCCTTCACCAGCCGCCAGCCTAGACAGCTACTCCACGTGCAATGTTTCCAGT 254
                                                                            26 AlaProAlaProProThrThr------PheGlyThrGluAspLeuValLeuThr 41
                                                                                                                                                                                                                      42 LeuAsnAsnGlnGlnMetThrLeuGluGly--------------
US-08-978-217-16_COPY_2_371 (1-370) x US-09-009-913-2 (1-5427)
                                                                                                                                                                                                                                                                                                                                                                    52 -----ProGluLysAlaSerTrpThrSerGlu-----
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qq	
Š	71 LeuGluTrplleSerTyrGlnValGluLygABnLysTyrAspAlaSerSer1leAspPhe 90
e e	315 TGGGAGTGGCTCCAGGACCTCCTGGACACCAGCTGGATGCCAATTGTATCCCTTTC 374
È	91 SerArgCysAsnMetAspGlyAlaThrLeuCysSerCysAlaLeuGluGluLeuArgLeu 110
Dp	375 CAAGAGTICGACATCAACGGCGAGCACCICIGCAGCAIGAGTITGCAGGAGTICACCCGG 434
ò	111 ValPheGlyProLeuGlyAspGlnLeuHisAlaGlnLeuArgAspLeuThrSerAsn 129
Dp.	435 GCGGCAGGAGCGCGGGGCAGCTCCTCTACAGCAACTTGCAGCATCTGAAGTGGAACGGC 494
₹	130SerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLyBAspGlyMetSer 147
q _Q	495 CAGTGCAGTAGTGACCTG 512
ζ.	148 PheGinGluSerLeuGlyAspLeuGlyProPheAspGlnGlySerProPheAlaGlnGlu 167
q _C	513 TTCCAGTCCACAATGTCATTGTCAAGACTGAACAAACTGAGCTTCC 563
ç,	168 LeuLeuAspAspGlyArgGlnAlaSerProTyrTyrCysSerThrTyrGlyFroGlyAla 187
do G	564 ATCATGAACACCTGGAAAGACGAGAACTATTATATGACACCAACTATGGT 614
δ,	188 ProSerProGlySerSerAspValSerThrAlaArgThrAlaThrProGlnSerSerHis 207
Db dc	615 620
Š	208 AlaSerAspSerGlyGlySerAspValAspLeuAspLeuThrGluSerLysValPhePro 227
QQ	621
λŏ	228 ArgAspAspPheThrAspTyrLysLysLysGlyGluProLysHisGly 242
qq	
ò	243 LysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCysLeuGluGly 262
qa	711 AAAAAGGAGCAAGACCCCCCTGCCAAGTGCCACAAA
8	263 LysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheIleArgAspIleLeu 282
අු	750AAGCACAACCCGAGAGGGACTCACTTATGGGAATTCATCCGCGACATCCTC 800
ò	283 IleHisProGlubeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGluGlyValPhe 302
qq	801 TTGAACCCAGACAAGAACCCAGGATTAATAAAATGGGAAGACCGATCTGAGGCGTCTTC 860
à	303 LysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLysAsnSerAsn 322
Op QC	::: :::
 70	323 MetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGlulleLeuGlu 342
qq	921 ATGACCTATGAAAAGCTCAGCCGAGCTATGAGATATTACTACAAAAGAGAAATACTGGAG 980
à	343 ArgValAspGlyArgArgLeuValTyrLysPheGlyLysAsnSerSerGlyTrpLysGlu 362
	981 CGTGTGGATGGACGAAGACTGGTATATAAATTTGGGAAGAATGCCCGAGGATGGAGAAA 1040
8	363 Gluglu 364
Db 1	041 AATGAA 1046
RESULT 10	

Sequence 3, Application US/09009913 Patent No. 6087485

US-09-009-913-3

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536 CAGCTCCTCTACAGCAACTTGCAGCATCTGAAGTGGAACGGCCAGTGCAGTAGTGAC--- 592
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               117 AspGlnLeuHisAlaGlnLeuArgAspLeuThrSerAsn-----SerSerAspGlu 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          134 LeuSerTrpIleIleGluLeuLeuGluLysAspGlyMetSerPheGlnGluSerLeuGly 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               593 -----CTGTTCCACACACACT 613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              154 AspleuGlyProPheAspGlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyArg 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     57 TrpThrSerGluArgProGlnPheTrpSerLysThrGlnValLeuGluTrpIleSerTyr 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 77 GlnValGluLysAsnLysTyrAspAlaSerSerileAspPheSerArgCysAsnMetAsp 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-978-217-16_COPY_2_371 (1-370) x US-09-009-913-3 (1-5510)
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Matches:
Conservative:
Mismatches:
Indels:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskethe
COMPUTER: Diskethe
COMPUTER: Diskethe
COMPUTER: Diskethe
COMPUTER: OF WINDOWS Version 2.0
SOFTWARE: PastEND for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
FILING DATE: 21-JAN-1998
CLASSIPICATION:
PRIOR APPLICATION:
APPLICATION NUMBER:
              APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
STATE: CA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,677
REPERENCE/DOCKET NUMBER: SEQ-4P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3331
TELEFAX: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS: LENGTH: 5510 base pairs TYPE: mucleic acid STRANDEDNES: double
                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
ATTORNEY/AGENT INFORMATION:
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54.43%
40.82%
29.29%
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Best Local Similarity:
GENERAL INFORMATION:
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TELEPHONE:

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                                                                                                                                                                                                                                                                                                                                                                                                  ProArgGlyThrHisLeuTrpGluPheIleArgAspIleLeuIleHisProGluLeuAsn 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       902 CCAGGATTAATAAAATGGGAAGACGGATCTGAGGGGGGTCTTCAGGTTCTTGAAATCAGAG 961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              309 AlaValAlaGlnLeuTrpGlyGlnLysLysLysAsnSerAsnMetThrTyrGluLysLeu 328
  174 GlnAlaSerProTyrTyrCysSerThrTyrGlyProGlyAlaProSerProGlySerSer 193
                                                                              194 AspValSerThrAlaArgThrAlaThrProGlnSerSerHisAlaSerAspSerGlyGly 213
                                                                                                                        703
                                                                                                                                                                                                                                          -----ThrAspTyrLysLysGlyGluProLysHisGlyLysArgLysArgGlyArg 248
                                                                                                                                                                                                                                                                                                                                                                                                                       704 ------GTAGATTTGTTGGACAGCAAAACTTTCTGCCGGGCTCAGATCTCCAIG 751
                                                                                                                                                                                                                                                                           752 ACAACCACCAGTCACCTTCCTGTTGCAGAGTCACCTGATATGAAAAGGAGCAAGACCC 811
                                                                                                                                                                                                                                                                                                                                                         812 CCTGCCAAGTGCCACACCAAA------------AAGCACAAC 841
                                                                                                                                                          214 SerAspValAspLeuAspLeuThrGluSerLysValPheProArgAspAspPhe----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     962 GCAGTGGCTCAGCTATGGGGTAAAAAGAAGAACAACAGCAGCAGCATGACCTTGATGAAAAAGCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      349 LeuvalTyrLysPheGlyLysAsnSerSerGlyTrpLysGluGluGlu 364
                             665 GACGAGAACTATTTATATGACACCAACTATGGT------
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEC for Windows Version 2.0
CURRENT APPLICATION DATA:
FILING DATE: 21-JAN-1998
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GENERAL INFORMATION:
APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
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STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
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                                                                                                                      -----AGCACA-----
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Patent No. 6087485
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TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
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999 CCGAGAGGGACTCACTTATGGGAATTCATCCGCGACACTCTCTTGAACCCAGACAAGAAC 1058
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               117 AspGlnLeuHisAlaGlnLeuArgAspLeuThrSerAsn-----SerSerAspGlu 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   134 LeuSerTrpllelleGluLeuGluLysAspGlyMetSerPheGlnGluSerLeuGly 153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             174 GlnAlaSerProTyrTyrCysSerThrTyrGlyProGlyAlaProSerProGlySerSer 193
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Conservative:
Mismatches:
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                                                                                                                                                                                                                                                                  Length:
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                                     INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 5667 base pairs
TYPE: nucleic acid
STRANDENESS: double
TOPOLOGY: linear
MOLECULE TYPE: CDNA
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                                                                                                                                                                                                                                                                                  580.00
54.43%
40.82%
29.29%
TELEFAX: 650-327-3231
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Best Local Similarity:
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COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO
       109 ArgleuValPheGlyProLeuGlyAspGlnLeuHisAlaGlnLeuArgAspLeuThrSer 128
                                         730 ACCCGGGCGGCAGGAGCGCGGGGGCANCTCCTCTACAGCAACTTGCAGCATCTGAAGTGG 671
                                                                                                    129 Asn-----SerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGly 145
                                                                                                                                                                                           146 MetSerPheGlnGluSerLeuGlyAspLeuGlyProPheAspGlnGlySerProPheAla 165
                                                                                                                                                                                                                                                                                      166 GlnGluLeuLeuAspAspGlyArgGlnAlaSerProTyrTyrCysSerThrTyrGlyPro 185
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GENERAL NO. GENERAL
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF
NUMBER OF SEQUENCES: 224
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
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; Sequence 44, Application US/09030607
; Patent No. 6262245
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COUNTRY: USI
ZIP: 98104
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APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Dillin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR INMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO
NUMBER OF SEQUENCES: 178
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                               309 AlavalAlaGlnLeuTrpGlyGlnLysLysLysAsnSerAsnMetThrTyrGluLysLeu 328
                                                                                            SerArgAlaMetArgTyrTyrLysArgGluIleLeuGluArgValAspGlyArgArg 348
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NOMBER: US/09/020,956
FILING DATE: 09-FEB-1998
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Indels:
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Patent No. 6261562
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NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 21012
REFERENCE/DOCKET NUMBER: 21012
TELECOMMUNICATION INFORMATION:
TELEFAX: (206) 622-4900
TELEFAX: (206) 622-6911
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 852 base pairs
TYPE: mucleic acid
TYPE: mucleic acid
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241 HisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCysLeu 260
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                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30
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Mismatches:
                                                                                                                                                   ATTORNEY AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.427C3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-690
TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 852 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length:
Matches:
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                                                                                                     APPLICATION NUMBER: US/09/030,607
FILING DATE: 25-FEB-1998
CLASSIFICATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Best Local Similarity:
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281 ileleullehisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGluGly 300
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261 GluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArgAsp
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APPLICANT: Red, Seven G.
APPLICANT: Ralos, Michael
APPLICANT: Ralos, Michael
APPLICANT: Ralos, Michael
APPLICANT: Retter, Mark
APPLICANT: Retter, Mark
APPLICANT: Day, Craig
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121-42709
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTUM NO 44
TEMBER OF SEQ ID NOS: 575
SOFTUM NO 44
TEMBER OF SEQ ID NOS: 575
TEMBER OF SEQ ID NOS: 575
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Mismatches:
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan Louise
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THER INFORMATION: n = A,T,C or G
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NAME/KEY: misc_feature
LOCATION: (1)...(852)
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Query Match:
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US-09-439-313-44/c
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TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
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Mismatches:
Indels:
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                 FILE REFERENCE: 210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
NUMBER OF SEQ ID NOS: 472
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 44
LENGTH: 852
                                                                                                                                                                                              ; OTHER INFORMATION: n = A,T,C or G
US-09-352-616A-44
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543.00
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40.79%
27.42%
                                                                                                                                                  FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(852)
                                                                                                                        TYPE: DNA
ORGANISM: Homo sapien
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Query Match:
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                                             109 ArgleuValPheGlyProLeuGlyAspGlnLeuHisAlaGlnLeuArgAspLeuThrSer 128
                                                                                                                                            129 Asn-----SerSerAspGluLeuSerTrplleIleGluLeuLeuGluLysAspGly 145
146 MetSerPheGlnGluSerLeuGlyAspLeuGlyProPheAspGlnGlySerProPheAla 165
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649 ---CTGTICCAGTCCACACATGTCATTGTCAAGACTGAACAAACTGAGCCT----
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Sequence 44, Application US/09352616A

Sequence 46, Application US/09352616A

Sequence 46, Application US/09352616A

GENERAL INFORMATION:
APPLICANT: Dillon, Davin C.
APPLICANT: Jang, Yuqui
APPLICANT: Xi, Jangchun
APPLICANT: Wi, Jangchun
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
                                                                                                                                                                          670 AACGGCCAGTGCAGTAGTGAC------
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                                                                        69 GlnValLeuGluTrpIleSerTyrGlnValGluLyBAsnLysTyrAspAlaSerSerIle 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    598 ---TCCATCATGAACACCTGGAAAGACNAGAACTATTTATATGACCAACTATGGT---
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US-08-978-217-16_COPY_2_371 (1-370) x US-09-352-616A-44 (1-852)
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CTGGAGCGTGTGGATGGACGAAGACTGGTATATAAATTTGGGAAGAAGAAGGATGGGATGG	CTGGAGCGTGTGATGGACGAAGACTGGTAT	184	a
341 LeuGluArgValAspGlyArgArgLeuValTyrLysPheGlyLysAsnSerSerGlyTrp 360	LeuGluArgValAspGlyArgArgLeuValT	341	ò
244 AGCAGCATGACCTATGAAAAGCTCAGCCGAGCTATGAGATATTACTACAAAAAGAGAAAATT 185	AGCAGCATGACCTATGAAAAGCTCAGCCGAG	244	q
321 SerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrTysLysArgGluIle 340	SerAsnMetThrTyrGluLysLeuSerArgA	321	Š
304 GTCTTCAGGTTCTTGAAATCAGAGGCAGTGGCTCAGCTATGGGGTAAAAAGAAGAACAAC 245	GTCTTCAGGTTCTTGAAATCAGAGGCAGTGG	304	q
301 ValPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysAsan 320	ValPheLysPheLeuArgSerGluAlaValA	301	ò

Search completed: February 12, 2004, 21:50:15 Job time : 105.27 secs

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Sequence 27, Appl
Sequence 4, Appli
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                                                                    February 12, 2004, 21:46:16; Search time 445.897 Seconds (without alignments) 7673.534 Million cell updates/sec
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1 GGATCCTTCCAAGGCACTGA......CAGAGGGGTCTCCTGGATCC 7752
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3: / cgn2_6/ptodata/1/ina/6A_COMB.seq:*
5: / cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: / cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: / cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-433-938-282
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US-09-016-434-927
US-09-026-941-4
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US-08-781-891-209
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US-09-188-930-75
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7 2 6 5 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	sequence 5, Applisequence 5, Applisequence 17, Applisequence 5, Applisequence 1, Applisequence 17, Applisequence 17, Applisequence 17, Applisequence 17, Applisequence 11, Applisequence 182, Applisequence	Targets and Methods	1907; 1907; 10dels

7293

1334 AAGCTGATGTTTTGGTGTATTGTCAGCCATCGTCGTTGGACTCGGAGACTATGGCCTCGC 1393 7239 GAAGAACCTGTTGGGCGTGCCCTCTGC---AGTCTCCTCAAGTGCAGCCTTTGGCCTC--

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7179 GGACAGGCAGGCCTGACGGCCCCCTTAACATGGATGTTCCCTGTGTTGCTGTAGAGAG 7238

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ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION: 514
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ATTORNEY/AGENT INFORMATION:
NAME: William T. Han
REGISTRATION NUMBER: 34,3
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                    7342 --ICGATAACTCTTCCAGCTGTGATTCCAGTTCCCTCCCGTCCCAACATGGACTGCAAAT 7399
                                                                                                                               1454 AGTGTATCTCCTTTTATCTGGTGCCTCCTCAAACCCAGTCTCAGACACTTAAATGCAGAC 1513
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                                                                                                                                                                                                                                                                                                                                                                                                                                            7573 IGCTATA--AATAITCCAGGIGIATATAGAGAGCIATITITITITAAAGCATITICCCCICC 7630
                                          1394 CTCCCCACCCTCCTCTTGGAATTACAAGCCCTGGGGTTTGAAGCTGACTTTATAGCTGCA 1453
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1095 AGGTACTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACAAG 1154
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2.8%; Score 216; DB 4; Length 1907;
Best Local Similarity 66.7%; Pred. No. 2.3e-49;
Matches 456; Conservative 0; Mismatches 195; Indels 33; Gaps
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; OTHER INFORMATION: Human epithelial-restricted with serine box (BSX)
; OTHER INFORMATION: protein.
US-09-570-593-4
7294 --TOTOCOTOGCOCTOTIGGAATTACAAGCCCCGGGTTTGAACCAACTIGT-----
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APPLICANT: Kaufmann, Joerg
APPLICANT: Kaufmann, Joerg
APPLICANT: Kin, Hong
ITILE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
FILE REFERENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FREUSEQ FOR WINDOWS Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1750 CACTCCTCTCCCACAGAGTGCTGG 1773
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Patent No. 6566063
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1394 CICCCCACCCTCCTCTGGAATTACAAGCCCTGGGGTTTGAAGCTGACTTATAGCTGCA 1453
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                                                                          1334 AAGCTGATGTTTTGGTGTATTGTCAGCCATCGTCCTTGGACTCGGAGACTATGGCCTCGC 1393
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                                                                                                                                                             7239 GAAGAACCIGITGGGCGTGCCCTCTGC---AGTCTCCTCAAGTGCAGCCTTTGGCCTC--
                                                                                                                                                                                                                                                                                                                               7294 --TCTCCTCGCCCTCTTGGAATTACAAGCCCCGGGTTTGAACCAACTTGT-----
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Patent No. 5789200

GENERAL INFORMATION:
APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
ITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3

NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
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6999 AGGIATTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGACGGCTCGTCTACAAG 7058
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1413 CTCCCCACCCTCCTCTTGGAATTACAAGCCCTGGGGTTTGAAGCTGACTTTATAGCTGCA 1472
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PREAT NO. 6518237
APPLICANT: Yuqui, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DAMPOSITIONS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C3
                                                                                                                                                                                                                      Query Match 2.8%; Score 215.6; DB 1; Length 1920; Best.Local Similarity 65.6%; Pred. No. 3e-49; Matches 449; Conservative 0; Mismatches 204; Indels 31;
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                       : LENGTH: 1920
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: No
SEQUENCE CHARACTERISTICS
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6316 AACGAAGGCCTCATGAAGTGGGAGAAACCGGCACGAGGGTGTGTTCAAGTTTCTTCGCTCA 6375
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Best Local Similarity 88.9%; Pred. No. 2.9e-35;
Matches 177; Conservative 0; Mismatches 22; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                  Gaps
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APPLICANT: Jiang, Yuqiu
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Mitchen, Jennifer L.
APPLICANT: Mitchen, Jennifer L.
APPLICANT: Mitchen, Jennifer L.
APPLICANT: Mitchen, Jennifer L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210111.47008
CURRENT APPLICATION NUMBER: US/09/620,405B
CURRENT APPLICATION NUMBER: 2000-07-20
NUMBER OF SEQ ID NOS: 495
SOFTWARE: FREESEQ for Windows Version 3.0
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CURRENT APPLICATION NUMBER: US/09/389,681A CURRENT FILING DATE: 1999-09-02 NUMBER OF SEQ ID NOS: 463 SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 282 LENGTH: 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 282, Application US/09620405B Patent No. 6528054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6436 CTGAGCCGAGCCATGAGGT 6454
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                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
US-09-389-681-282
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CRGANISM: Homo sapiens
US-09-620-4058-282
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LENGTH: 502
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ORGANISM: Homo sapiens
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US-09-604-287A-282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               258 GCGCCCAGAGGCACCCACCTGTGGGAGTTCATCCGGGACATCCTCATCCACCCGGAGCTC 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            318 AACGAGGCCTCATGAAGTGGGAGAATCGGCATGAAGGCGTCTTCAAGTTCCTGCGCTCC 377
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitchai, Jennifer L.
APPLICANT: Mitchai, Jennifer L.
APPLICANT: Mitchai, Jennifer L.
APPLICANT: Mitchai, Jennifer L.
APPLICANT: Mit Jiangchun
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DOMPOSITIONS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C4
CURRENT FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 474
SOSTWARE: FARELSEQ for Windows Version 3.0
                                                                                                                     Sequence 282, Application US/09339338A

Sequence 282, Application US/09339338A

Patent No. 6573368

GENERAL INFORMATION:
APPLICANT: Vigitu, Jiang

APPLICANT: Witcham, Janifer L.
APPLICANT: Mitcham, Janifer L.
APPLICANT: No. Jiangchun

TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.47002

CURRENT APPLICATION NUMBER: US/09/339,338A

CURRENT FILING DATE: 1999-06-23

NUMBER OF SEQ ID NOS: 315

SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.1%; Score 163.8; DB 4; Length 502;
88.9%; Pred. No. 2.9e-35;
tive 0; Mismatches 22; Indels 0
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Patent No. 6579973
GENERAL INFORMATION:
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438 CTGAGCCGGCCATGAGGT 456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens
US-09-339-338-282
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
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US-09-433-826B-282
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                                                                           RESULT 6
US-09-339-338-282
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6376 GAGGCCGTGGCCCAACTCTGGGGCCAGAAGAAGAAGAACAGCAACATGACCTATGAGAAG 6435
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                                                                                                                                                                                                                                                                                                         6376 GAGGCCGTGGCCCAACTCTGGGGCCAAGAAGAAGAAGAACAGCAACATGACCTATGAGAAG 6435
6256 GCCCCCAGAGGTACTCACCTGTGGGAGTTTATCCGAGACATCCTAATCCACCCCGAGCTC 6315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           378 GAGGCTGTGGCCCGAACTATGGGCCCAAAAGAAAAGAACAGCAACATGACCTACGAGAAG 437
                                                                            258 GCGCCAGAGAGGCACCCACCTGTGGGAGTTCATCCGGGACATCCTCATCCACCCGGAGCTC 317
                                                                                                                                                                                                                         318 AACGAGGGCCTCATGAAGTGGGAAATCGGCATGAAGGCGTCTTCAAGTTCCTGCGCTCC 377
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Miccham, Jennifer L.
APPLICANT: Miccham, Jennifer L.
APPLICANT: Ku, Jiangchun
APPLICANT: Ku, Jiangchun
APPLICANT: Ku, Jiangchun
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.47007
CURRENT FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING TITLE OF INVENTION: PATHWAY GENE EXPRESSION NUMBER OF SEQUENCES: 1450
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 2.1%; Score 163.8; DB 4; Length 502; Best Local Similarity 88.9%; Pred. No. 2.9e-35; Matches 177; Conservative 0; Mismatches 22; Indels 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 282, Application US/09604287A Patent No. 6586572
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Patent No. 6500938
GENERAL INFORMATION:
APPLICANT: Janice Au-Young
APPLICANT: Jeffrey J. Seilhamer
                                                                                                                                                                                                                                                                                                                                                                                                                                                          6436 CTGAGCCGAGCCATGAGGT 6454
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Sequence 4, Application US/09245041

Sequence 4, Application US/09245041

Patent No. 6274339

GENERAL INFORMATION:
APPLICANT: Mocre, N.
APPLICANT: Nagle, D.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND TREATMENT TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY
FILE REPERENCE: 7883-136
CURRENT APPLICATION NUMBER: US/09/245,041
CURRENT FILING DATE: 1999-02-05
EARLIER APPLICATION NUMBER: 60/093,630
EARLIER FILING DATE: 1998-07-21
EARLIER APPLICATION NUMBER: 60/104,978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7059 TTTGGCAAGAACTCTAGTGGCTGGAAGGAAGAAGAGGTTGGAGAGAGTCGGAATTAAGGA 7118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6999 AGGIATIACIACAAACGGGAGAICCIGGAACGGGIGGAIGGAIGGCCGACGGCTCGICTACAAG 7058
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                                                                                                                                                    COMPUTE: 74204
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: EMP PC compatible
CORRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/016,434
FILLING DATE: HEREWITH
SSEE: INCYTE PHARMACEUTICALS, INC.
I: 3174 PORTER DRIVE
PALO ALTO
CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
NAME: Zeller, Karen J.
NEGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: 94-0002 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 845-4166
INFORMATION FOR SEQ ID NO: 927:
SEQUENCE CHARACTERISTICS:
LENGTH: 237 base pairs
TYPE: nucleic acid
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Best Local Similarity 79.3
Matches 149; Conservative
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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CLONE: 773734
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           ADDRESSEE:
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                                                                                                                     COUNTRY:
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US-09-020-956-44/c
US-09-020-956-44/c

J Sequence 44, Application US/09020956

J Patent No. 626156.

J APPLICANT: Xu, Jiangchun

APPLICANT: Xu, Jiangchun

APPLICANT: Xu, Jiangchun

APPLICANT: Nu, Jiangchun

TITLE OF INVENTION: Compounds FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO

NUMBER OF SEQUENCES: 178

CORRESPONDENCE ADDRESS:

ADDRESSEE: SEED and BERRY LLP

STREET: 6300 Columbia Center, 701 Fifth Avenue

CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                      1841 TITGITCTGTAGITTTTAGTTTTTCAAGACAGAGITTCTCTGTGTAGCCCTGGCTGTCCTG 1900
                                                                                                                                                                                                                                                                                                                                                                                                    5390 Trontuncantincecritingaacaeerricicieneraeeccieecre
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0
                                                                                                                                                                                                                                           Length 5973;
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1.6%; Score 123.6; DB 3; Length !
Best Local Similarity 67.4%; Pred. No. 2e-23;
Matches 174; Conservative 0; Mismatches 84; Indels
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/020,956
FILING DATE: 09-FEB-1998
CLASSIFICATION:
ATTORNEY/APPLICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.427C2
TELECOMMUNICATION INFORMATION:
                         NUMBER OF SEQ ID NOS: 131
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2081 TGTTGGGCTTCTGACCAC 2098
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EARLIER FILING DATE: 1998-10-20
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TELEPAX: (206) 628-6031
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 852 base pairs
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                              TYPE: DNA ORGANISM: Mus musculus
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                                                                          SEQ ID NO 4
LENGTH: 5973
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MOLECULE TYPE:

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6378 GGCCGTGGCCCAACTCTGGGCCCAGAAGAAGAACAGCAACATGACCTATGAGAAGCT 6437
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                                                                       6318 CGAAGGCCTCATGAAGTGGGAAACCGGCACGAGGGTGTGTTCAAGTTTCTTCGCTCAGA 6377
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                                                                                                                                                                                                             281 GGCAGTGGCTCAGCTATGGGGTAAAAAGAAGAACAACAACAGCATGACTGATGAAAAGCT 222
                       401 cccgagagagagrenerirangagarinearccagacarccicricaacccagacaagaa 342
                                                                                                                      341 cccaddarraaraaraddaaagacdarcrdaddddcdrcrrcaddrrcrrgaaarcada 282
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APPLICANT: Read, Steven G.

APPLICANT: Rades, Michael
APPLICANT: Rades, Michael
APPLICANT: Reflex, Mark
APPLICANT: Setter, Mark
APPLICANT: Solk, John
APPLICANT: Solk, John
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FREASE STS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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74.1%; Pred. No. 1.1e-21;
tive 0; Mismatches 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 14
US-09-352-616A-44/c
; Sequence 44, Application US/09352616A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang Yuqui
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 44, Application US/09439313; Patent No. 6329505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , LOCATION: (1)...(852)
, OTHER INFORMATION: n = A,T,C or G
US-09-439-313-44
                                                                                                                                                                                                                                                                        6438 GAGCCGAGCCATGAGGT 6454
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Best Local Similarity 74.14
Matches 146; Conservative
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Sequence 44, Application US/09030607
Patent No. 626245
GENERAL INFORMATION:
APPLICANT: Mu. Jiangchun
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO
NUMBER OF SEQUENCES: 224
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
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                                                                                                                                                                                                6258 CCCCAGAGGTACTCACCTGTGGGAGTTTATCCGAGACATCCTAATCCACCCCGAGCTCAA 6317
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                                                                                               Query Match
1.5%; Score 115.4; DB 3; Length 852;
Best Local Similarity 74.1%; Pred. No. 1.1e-21;
Matches 146; Conservative 0; Mismatches 51; Indels 0
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1.5%; Score 115.4; DB 3; Length 852;
Best Local Similarity 74.1%; Pred. No. 1.1e-21;
Matches 146; Conservative 0; Mismatches 51; Indels 0
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: DEC compatible

OPERATING SYSTEM: PC-DOS/NS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/030,607

FILING DATE: 25-FEB-1998

CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           210121.42703
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NAME: Maki, David J.
REGISTATION NUMBER: 31,392
REPERBUCE/DOCKET NUMBER: 21012
TELECOMMUNICATION INFORMATION:
TELEFAX: (206) 622-4900
TELEFAX: (206) 622-631
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 852 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                       ORGANISM: Homo sapiens
US-09-020-956-44
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ORGANISM: Homo san
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ORIGINAL SOURCE:
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GENERAL INFORMATION:
APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Micham, Jennifer Lynn
APPLICANT: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OPPROSTATE CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION NUMBER: US/09/352,616A
CURRENT APPLICANTON NUMBER: US/09/352,616A
NUMBER OF SOI ID NOS: 472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 1.5%; Score 115.4; DB 4; Length 852; Best Local Similarity 74.1%; Pred. No. 1.1e-21; Matches 146; Conservative 0; Mismatches 51; Indels 0
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US-09-232-149A-44/C

Sequence 44, Application US/09232149A

Sequence 44, Application US/09232149A

Sequence 44, Application US/09232149A

Setent No. 6465611

GENERAL INFORMATION:
APPLICANT: Witcham, Januagehun

APPLICANT: Mitcham, Jennifer Lynn

APPLICANT: Mitcham, Jennifer Lynn

APPLICANT: Mitcham, Jennifer Lynn

TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE

FILE REPERENCE: 210121.427C6

CURRENT APPLICATION NUMBER: US/09/232,149A

CURRENT FILING DATE: 1999-01-15

NUMBER OF SEQ ID NOS: 338

SOFTWARER FESTERE FESTERE FOR Windows Version 3.0
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1.5%; Score 115.4; DB 4; Length 852;
Best Local Similarity 74.1%; Pred. No. 1.1e-21;
Matches 146; Conservative 0; Mismatches 51; Indels 0;
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LOCATION: (1)...(852)

CTHER INFORMATION: n = A,T,C or G

US-09-352-616A-44
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LCCATION: (852)

/ OTHER INFORMATION: n = A,T,C or G

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ORGANISM: Homo sapien
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Best Local Similarity
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SEQ ID NO 44
LENGTH: 852
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Sequence 927, App
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7673.534 Million cell updates/sec
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-620-105B-282

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US-09-630-958A-27

US-09-570-958A-27

US-09-570-958A-27

US-09-570-593-4

US-08-7479-728B-1

US-08-7479-728B-1

US-09-103-840A-1

US-09-103-840A-1

US-09-103-840A-1

US-09-103-840A-1

US-09-103-840A-1

US-09-103-840A-1

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US-09-495-050A-190
US-09-009-913-338
US-09-535-008-55
US-09-350-457A-1
US-08-352-902D-145
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Maximum Match 100%
Listing first 45 summaries
                                                                                               OM nucleic - nucleic search, using sw model
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21
1 GTACCTCATGGCCCGGCTCAG 21
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Gapop 10.0 , Gapext 1.0
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Match
                                                                                                                                                                                                                                               Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                    Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                              Searched:
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Sequence 145, App

Sequence

2 8 15.2 72.4 2484 3 US-08-291-8 2 9 15.2 72.4 2484 3 US-08-361-901-4 2 15.2 72.4 2484 4 US-09-961-901-901-901-901-901-901-901-901-901-90	Sequence 8, Appli Sequence 4, Appli Sequence 4, Appli Sequence 16, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 240, Appli Sequence 5, Appli Sequence 5, Appli Sequence 1, Appli Sequence 200, Appli	I GNAL I NG	237;
C 28 15.2 72.4 2484 C 30 15.2 72.4 2484 C 31 15.2 72.4 2484 C 32 15.2 72.4 2484 C 32 15.2 72.4 2484 C 33 15.2 72.4 2484 C 33 15.2 72.4 2525 C 34 15.2 72.4 2667 C 34 15.2 72.4 2667 C 34 15.2 72.4 2667 C 37 15.2 72.4 2667 C 38 15.2 72.4 2667 C 39 15.2 72.4 2667 C 39 15.2 72.4 4953 C 41 14.8 70.5 1030 C 42 11.8 70.5 1030 C 58QUENCE PROPARION: DATA CONPUTER: READABLE FORM: CONPUTER: BADO ALTO CONPUTER: DATO ALTO CONPUTER: DATO ALTO CONPUTER: READABLE FORM: CONPUTER: DATO ALTO CONPUTER: PADO ALTO CONPUTER: DATO ALTO CONPUTER:	US-08-209-521-8 US-08-961-810-4 US-08-355-902D-4 US-09-265-503B-4 US-08-200-16 US-08-200-16 US-08-468-024B-1 US-08-468-024B-1 US-08-468-024B-1 US-08-468-024B-1 US-09-620-121-1 US-09-620-312-240 US-09-333-599-5 US-09-333-599-5 US-09-333-599-1 US-09-333-599-1 US-09-333-599-1 US-09-333-599-1 US-09-333-599-1 US-09-333-599-1 US-09-333-599-1 US-09-398-416-20 US-09-38-117-5 US-09-095-117-5	HE DETECTION OF ESSION INC.	Score 21; DB 4; Pred. No. 0.3;
	28 29 31 31 31 32 33 34 35 36 37 37 38 38 39 39 40 41 41 41 41 42 44 44 44 44 44 44 44 44 44	RESULT 1 US-09-016-434-927/C US-09-016-434-927/C Sequence 927, Application I Patent No. 650039 GENERAL INFORMATION: APPLICANT: Jeffrey J. S TILLE OF INVENTION: PAI NUMBER OF INVENTION: PAI NUMBER OF INVENTION: PAI NUMBER OF SEGUENCES: J. STREET: J. PALO ALTO STREET: ALLO ALTO STREET: ALLO ALTO COMPUTER: IBM PC COMPUTER: IBM PC COMPUTER: Word Perfect CLASSIFICATION NUMBER: FILING DATE: FILING DATE: HERBWITT APPLICATION NUMBER: FILING DATE: CLASSIFICATION NUMBER: FILING DATE: ATTORNEY/ACBNT INFORMATI NAME: Zeller, Karen GREDIED NO: SEQUENCE CHARACTERISTICE LERGYMUNICATION NUMBER: TELEPHONE: GEOD 855-TELEPHONE: GEOD 855-TELEPHONE: GEOD 855-TELEPHONE: GEOD 855-TELEPHONE: GEOD 855-TELEPHONE: GEOD 855-TELEPHONE: GEOD 855-TENERHITE SOURCE: LIBRARY: COLNCRIOL SEQUENCE CHARACTERISTICE LERGYH: Z37 Dase Pail TYPE: ILBRARY: COLNCRIOL SEQUENCE: CLONERICE: LIBRARY: COLNCRIOL US-09-016-434-927	Match Local Similarity

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Query Match
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Sequence 282, Application US/09389681A

Sequence 282, Application US/09389681A

Sequence 282, Application US/09389681A

Sequence 282, Application

Sequence 282, Application

APPLICANT: Vuqui, Jiang

APPLICANT: Vuqui, Jiang

APPLICANT: Xi, Jiangchun

TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND

TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE

FILE REFERENCE: 210121.470C3

CURRENT APPLICATION NUMBER: US/09/389,681A

CURRENT FILING DATE: 1999-09-02

NUMBER OF SEQ ID NOS: 463

SOFTWARE: FREESE FREESE FOR Windows Version 3.0
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     Gaps
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Sequence 282, Application US/09620405B

Patent No. 628054

GENERAL INFORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: William T.
APPLICANT: William T.
APPLICANT: William T.
TITLE OF INVENTION: COMPOSITIONS OF BREAST CANCER
FILE REFERENCE: 210121.47008

CURRENT APPLICATION UNMBER: US/09/620,405B

CURRENT APPLICATION OF 2495

NUMBER OF SEQ ID NOS: 495
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  0; Mismatches
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                                               1 GTACCTCATGGCCCGGCTCAG 21
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21; Conservative
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US-09-389-681-282
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ORGANISM: Homo sapiens
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US-09-339-338-282/c

RESULT 4

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Sequence 282, Application US/0933938A
Patent No. 657356
GENERAL INFORMATION:
APPLICANT: Yugiu, Jiang
APPLICANT: Yugiu, Jiang
APPLICANT: Wigiu, Jiangchun C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Xu, Jiangchun C.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGROSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C2
CURRENT APPLICATION NUMBER: US/09/339,338A
CURRENT FILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 315
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 282
LENGTH: 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jiang, Yuqui
APPLICANT: Jiang, Yuqui
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangcham, Jennifer L.
APPLICANT: Xu, Jiangcham, Jennifer L.
APPLICANT: Xu, Jiangcham, Jennifer L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121-470C4
CURRENT APPLICATION NUMBER: US/09/433,826B
CURRENT APPLICATION NUMBER: US/09/433,826B
SUPTHMER OF SEQ ID NOS: 474
SOFTWARE: FastSEQ for Windows Version 3.0
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APPLICANT: Harlocker, Susan L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 21; DB 4; Length 502; 100.0%; Pred. No. 0.31;
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Best Local Similarity 100.0%; Score 21; DB 4; Length 502;
Best Local Similarity 100.0%; Pred. No. 0.31;
Matches 21; Conservative 0; Mismatches 0. Tradele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
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Sequence 282, Application US/09433826B

; Patent No. 657997;

; GENERAL INFORMATION:
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; Sequence 282, Application US/09604287A
; Patent No. 6586572
; GENERAL INFORMATION:
; APPLICANT: Diang, Yaqiu
; APPLICANT: Dillon, Davin C.
; APPLICANT: Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              458 GTACCTCATGGCCCGGCTCAG 438
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Matches 21; Conservative
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CORGANISM: Homo sapiens
US-09-339-338-282
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SOFTWARE: FastSEQ for Windows Version 4.0
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| Sequence 27, Application US/09300958A
| Patent No. 6495319 |
| GENERAL INFORMATION |
| APPLICANT: McClelland, Michael |
| APPLICANT: Welsh, John |
| APPLICANT: Welsh, John |
| APPLICANT: TERKIA, Thomas |
| TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of TITLE OF INVENTION: Bing Same |
| TITLE OF INVENTION: Using Same |
| FILE REPRENCE: P-PH 3457 |
| CURRENT FILING DATE: 1999-04-27 |
| PRIOR FILING DATE: 1998-04-27 |
| PRIOR FILING DATE: 1998-03-27 |
| PRIOR FILING DATE: 1998-03-27 |
| PRIOR FILING DATE: 1998-02-04 |
| PRIOR FILING DATE: 1999-02-04 |
| PRIOR FILING DATE: 1998-03-27 |
| PRIOR FILING DATE: 1998-03-04 |
| PRIOR FILING DATE: 1998-04
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APPLICANT: Kaufmann, Joerg
APPLICANT: Kin, Hong
APPLICANT: Ain, Hong
APPLICANT: Ain, Hong
APPLICANT: Ain, Hong
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
FILE REFERENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER FILE REFERENCE: 210121.470C7
CURRENT PAPLICATION NUMBER: US/09/604,287A
CURRENT FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 22
LENGTH: 502
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Patent No. 6566063
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Matches 21, Conservative
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Best Local Similarity 100.
Matches 21; Conservative
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                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-604-287A-282
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LENGTH: 1907
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                                                                                  TYPE: DNA
CREANISM: Homo sapiens
CREANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (96)...(1211)
OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)
COTHER INFORMATION: protein.
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Patent No. 5789200
GENERAL INFORMATION:
APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
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100.0%; Score 21; DB 4; Length 1907;
Best Local Similarity 100.0%; Pred. No. 0.33;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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100.0%; Score 21; DB 1; Length 1920;
Best Local Similarity 100.0%; Pred. No. 0.33;
Matches 21; Conservative 0; Mismatches 0; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
COMPUTER: IBM 486
COMPUTER: IBM 406
COMPU
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ATTORNEY/AGENT INFORMATION:
NAME: William T. Han
REGISTRATION NUMBER: 34,344
REFREENCE/DOCKET NUMBER: ATG
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270 5219
TELEFAX: 610 270 4026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: Linear ANTI-SENSE: NO
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SEQ ID NO 4
LENGTH: 1907
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ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433
            STREET: F.C.
                                                                      Texas
                                                                              COUNTRY: Un
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| Sequence 17, Application PC/TUS9502251
| GENERAL INFORMATION: GENERAL INFORMATION: TITLE OF INVENTION: TITLE OF INVENTION: CELLS
| TITLE OF INVENTION: CELLS
| NUMBER OF SEQUENCES: 18
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 78.1%; Score 16.4; DB 3; Length 5499; Best Local Similarity 94.4%; Pred. No. 49; Matches 17; Conservative 0; Mismatches 1; Indels 0.
                                                                                GENERAL INFORMATION:
APPLICANT: Bonadio, Jeffrey
APPLICANT: Yin, Wushen
TITLE OF INVENTION: LATENT TGF (BINDING PROTEIN (LTBP)
TITLE OF INVENTION: GENES, COMPOSITIONS AND METHODS
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSES: Williams, Morgan & Amerson
STREET: 7676 Hillmont, Suite 250
                                                                                                                                                                                                                                                                                                                                                                        COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEB FC compatible
COMPUTER: TEB FC compatible
COMPUTER: TEB FC compatible
COMPUTER: TEB FC compatible
COMPUTER: TEB FC compatible
COMPUTER: PatentIn Release #1.0, Version #1.30
CURSTRING SYSTEM: DG-DG/N8-DGS
CLASSIFICATION NUMBER: US/08/479,722B
FILING DATE: 0.7-JUN-1995
CLASSIFICATION NUMBER: US PCT/US95/02251
FILING DATE: 21-FEB-1995
FILING DATE: 30-SEP-1994
FILING DATE: 18-FEB-1994
FILING DATE: 18-FEB-1994
APPLICATION NUMBER: US 08/199,780
FILING DATE: 18-FEB-1994
ATTORNEY AGENT INFORMATION:
NAMME: FUSSEY, Shelley P.M.
REGISTRATION NUMBER: 319,458
REFERENCE/DOCKET NUMBER: 4100.000500/FUS
TELLECOMMUNICATION INDER: 310.000500/FUS
TELLECOMMUNICATION INDER: 310.000500/FUS
RESULT 10
1S-08-479-722B-1/c
1 Sequence 1, Application US/08479722B
Patent No. 6074840
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (713) 934-7000
TELEPAX: (713) 934-7011
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 5499 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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; LOCATION: 1..5499
US-08-479-722B-1
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                                                                                                                                                                                                                                                                                                                     Texas
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US-09-213-767-1/c
; Sequence 1, Application US/09213767
; Batent No. 5448680
; GENERAL INFORMATION:
    APPLICANT: Lex M. COWSET
; TITLE OF INVENTION: ANTISENSE MODULATION OF ELK-1 EXPRESSION
    TITLE OF INVENTION: ANTISENSE WOULATION OF ELK-1
; CURRENT APPLICATION UNMERS: US/09/213,767
; CURRENT FILING DATE: 1998-12-17
; NUMBER OF SEQ ID NOS: 47
; ERNGTH: 2266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 78.1%; Score 16.4; DB 5; Length 5502; Best Local Similarity 94.4%; Pred. No. 49; Matches 17; Conservative 0; Mismatches 1; Indels 0
                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC COMPATIBLE
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: Parentin Release #1.0, Version
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/USS5/02251
FILING DATE: 30.5EP-1994
CLASSIFICATION
APPLICATION NUMBER: US 08/199,780
FILING DATE: 18 FEB-1994
CLASSIFICATION
APPLE: 18 FEB-1994
CLASSIFICATION:
ANABE: PARKET, DAVIG L.
REGISTRATION NUMBER: 32,165
REERENCE/DOCKET NUMBER: UNICOO9P--
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELEFAX: (312) 789-2679
United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
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INPORMATION POR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 5502 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
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US-09-213-767-1
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; NAME/KEY: CDS
; LOCATION: 1..5502
PCT-US95-02251-17
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Search completed: February 13, 2004, 01:55:01 Job time : 20.2079 secs
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US-09-103-840A-1
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OTHER INFORMATION: "n" bases at various positions throughout the sequence of CHER INFORMATION: represent a, t, c or g
US-09-103-840A-2
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OTHER INFORMATION: "n" bases at various positions throughout the sequence
OTHER INFORMATION: represent a, t, c or g
                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: FILEISCHMAN, Robert D.
APPLICANT: WHITE, Owen R.
APPLICANT: WHITE, Owen R.
APPLICANT: WHITE, Owen R.
APPLICANT: WINTER, John C.
ITILE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM TITLE OF INVENTION: TUBRECULOSIS
FILE REFERENCE: 24366-20007.00
GURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: WHITE, Owen R.
APPLICANT: WHITE, Owen R.
APPLICANT: PRASER, Claire M.
APPLICANT: PRASER, Claire M.
APPLICANT: PRASER, Claire M.
TITLE OF INVENTION: TUBERCULOSIS
TITLE OF INVENTION: TUBERCULOSIS
FILE REPERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LENGTH: 4403765
                                                   Gaps
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Query Match 77.1%; Score 16.2; DB 2; Length 2266; Best Local Similarity 85.7%; Pred. No. 59; Matches 18; Conservative 0; Mismatches 3; Indels 0;
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                                                                                                                                           513 GTACCGCAAGGCCCGGCTGAG 493
                                                                                             1 GTACCTCATGGCCCGGCTCAG 21
                                                                                                                                                                                                                                                                  Sequence 2, Application US/09103840A Patent No. 6294328
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION: APPLICANT: FLEISCHMAN, Robert D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             77.18;
85.78;
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Best Local Similarity
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LENGTH: 4403765
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US-09-103-840A-2/c
                                                                                                                                                                                                                   RESULT 13
US-09-103-840A-2
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APPLICANT: FRASER, Claire M.
APPLICANT: FRASER, Claire M.
APPLICANT: FRASER, Claire M.
APPLICANT: WENTER, Obn C.
TITLE OF INVENTION: TUBERCULOSIS
FILE REFRENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PATENTIN Ver. 2.1
  0; Gaps
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85.7%; Pred. No. 54;
tive 0; Mismatches 3; Indels 0;
3; Indels
  0; Mismatches
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ORGANISM: Mycobacterium tuberculosis

OTHER INFORMATION: H37Rv
US-09-103-840A-1
                                                                                                                                                                                                                  ; Sequence 1, Application US/09103840A; Patent No. 2594328
; BERERAL INFORMATION:
; APPLICANT: FLEISCHMAN, Robert D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 85.74
Matches 18; Conservative
  Matches 18; Conservative
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1002
1006
1006
11004
11004
11320
11320
11320
11320
11515
11515
11527
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US-09-389-681-282
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LENGTH: 502
         TYPE: DNA
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303, App
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Sequence 282, App
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                                                         February 12, 2004, 21:46:16; Search time 1.20792 Seconds (without alignments) 7673.534 Million cell updates/sec
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Sequence
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
       GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-252-991A-10598
US-09-620-312D-287
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US-09-433-826B-282
US-09-604-287A-282
US-09-570-593-4
US-08-746-789A-1
US-09-328-359-9
US-09-328-789-1
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                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
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-09-643-597-303
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US-08-826-246-9
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-08-925-588-9
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                                                                                                                                                                  569978 segs, 220691566 residues
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Maximum Match 100%
Listing first 45 summaries
                                          - nucleic search, using sw model
                                                                                                                    1 CCGGGACATCCTCATCCACCC 21
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                                                                                                                                                                                                       Minimum DB seq length: 0
Maximum DB seq length: 200000000
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21
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Match Length DB
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1986
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                 880, App
4285, Ap
1668, Ap
11668, Ap
11872, A
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8, Appli
11508, A
5, Appli
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APPLICANT: Yuqui, Jiang
APPLICANT: Yuqui, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: LIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210121.470C3
CURRENT APPLICATION NUMBER: US/09/389, 681A
CURRENT FILING DATE: 1999-09-02
NUMBER OF SEQ ID NOS: 463
SOFTWARE: FREESEQ for Windows Version 3.0
10256,
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; Sequence 282, Application US/09620405B
; Patent No. 6528054
; GENERAL INFORMATION:
; APPLICANT: Dillon, Davin C.
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Harlocker, Susan L.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
; FILE REPERENCE: 210121.470c8
; CURRENT FILING DATE: 2000-07-20
; NUMBER OF SEQ ID NOS: 495
; SOFTWARE: FRANCEQ for Windows Version 3.0
; SEQ ID NO 282
                                                           Sequence
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100.0%; Pred. No. 0.92;
tive 0; Mismatches 0; Indels
                                                                                            US-09-252-991A-11872
US-09-129-033-1
US-09-516-914-15054
US-09-516-914-16
US-09-592-074-1
US-09-252-91A-856
US-08-461-775-8
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US-09-252-991A-11508
                                    US-09-252-991A-4285
US-09-702-705-1668
US-09-736-457-1668
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US-09-252-991A-815
US-08-461-775-10
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US-09-031-606-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 282, Application US/09389681A; Patent No. 6518237; GENERAL INFORMATION:
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Matches 21, Conservative
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILLE REFERENCE: 21011.47004
CURRENT APPLICATION NUMBER: US/09/433,826B
CURRENT FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 474
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 282, Application US/09339338A

Batent No. 6573368

General Information US/09339338A

APPLICANT: Vigiu, Jiang

APPLICANT: Micham, Davin C.

APPLICANT: Mi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
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Thes 0; Indels
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Dillon, Davin C.
Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
CORGANISM: Homo sapiens
US-09-433-826B-282
                                           ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-620-405B-282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Homo sapiens
US-09-339-338-282
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APPLICANT: Jiang.
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US-09-339-338-282
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LENGIH: 502
LENGTH: 502
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LENGTH: 502
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1 CCGGGACATCCTCATCCACCC 21

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GENERAL INFORMATION:
APPLICANT: McClelland, Michael
APPLICANT: Welsh, John
APPLICANT: Tenkle, Thomas
TITLE OF INVENTION: Using Same
FILE REFERENCE: P-PH 3457
CURRENT APPLICATION NUMBER: US/09/300,958A
CURRENT APPLICATION NUMBER: 06/083,331
PRIOR APPLICATION NUMBER: 60/083,331
PRIOR FILING DATE: 1998-04-27
PRIOR FILING DATE: 1998-04-27
PRIOR FILING DATE: 1998-08-27
PRIOR PLICANION NUMBER: 60/098,070
PRIOR FILING DATE: 1998-08-27
PRIOR APPLICANION NUMBER: 60/118,624
PRIOR PLILING DATE: 1999-02-04
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                                                                                                                                                                                                                         APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: AL, Jiangchun
APPLICANT: Harlocker, Susan L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REPERENCE: 210121.47007
CURRENT APPLICATION NUMBER: US/09/604,287A
CURRENT APPLICATION NUMBER: US/09/604,287A
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FREESQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                             ; Sequence 282, Application US/09604287A; Patent No. 6586572
290 CCGGGACATCCTCATCCACCC 310
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; Sequence 27, Application US/09300958A
; Patent No. 6495319
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 27
LENGTH: 1907
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Best Local Similarity 100.0
--- Abes 21, Conservative
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US-09-604-287A-282
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; ORGANISM: Homo sapiens
US-09-300-958A-27
                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                 RESULT 5
US-09-604-287A-282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 282
LENGTH: 502
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; LOCATION: (96)...(1211)

; OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)

; OTHER INFORMATION: protein.

US-09-570-593-4
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Patent No. 5789200

GENERAL INFORMATION:

APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
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100.0%; Score 21; DB 4; Length 1907;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 21; Conservative 0; Mismatches 0; Indels (
                                                                                              APPLICANT: Kaufmann, Joerg
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Win, Hong
APPLICANT: Harrowe, GREESION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
FILLE REPRENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT PILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: PA
COUNTRY: USA
ZIP: 19406.04
ZIP: 19406.04
ZIP: 19406.04
ZIP: 19406.04
ZIP: 19406.04
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746,789A
FILING DATE: NO. 57892006mber 15, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
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REGISTRATION NUMBER: 34,344
REFERENCE/DOCKET NUMBER: ATG
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270 5219
TELEPHONE: 610 270 4026
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                 // Sequence 4, Application US/09570593
// Patent No. 6566063
// GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 4
LENGTH: 1907
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US-08-746-789A-1
US-09-570-593-4
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US-09-328-352-99/C
US-09-328-352-99/C
US-09-328-352-99/C
US-09-328-352-99/C
Sequence 99, Application US/09328352
Patent No. 6562958
GENERAL INFORMATION:
APPLICANT: GATY L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNI FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: BAUMANNI FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT PILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
LENGTH: 1086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
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Pred. No. 1.1e+02;
0; Mismatches 1; Indels 0;
                                                                                                                                                                 Query Match 100.0%; Score 21; DB 1; Length 1920; Best Local Similarity 100.0%; Pred. No. 1; Aatches 21; Conservative 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Lemke Ph.D. et al., Greg E.
TITLE OF INVENTION: PROTEIN-TYROSINE KINASE GENES
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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CLASSIPICATION: 530
PRIOR APPLICATION NUMBER: US 08/237,401
FILING DATE: 02-MAY-1994
PRIOR APPLICATION DATE: US 08/237,401
FILING DATE: 15-MAY-1994
APPLICATION NUMBER: US 07/884,486
FILING DATE: 15-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/456,647B
FILING DATE: 02-UN-1995
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM in MEDIUM TYPE: Floppy disk CMPUTER: IEM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                               951 ccededacarcercarceacec 971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/08456647B
Patent No. 5811516
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                   1 CCGGGACATCCTCATCCACCC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) ORGANISM: Acinetobacter baumannii
US-09-328-352-99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 94.4%;
Matches 17; Conservative
                    TYPE: Nucleic Acid
STANDEDNESS: Single
TOPOLOGY: Linear
JANTI-SENSE: NO
US-08-746-789A-1
LENGTH: 1920
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US-08-456-647B-1/c
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US-09-643-597-303
                                   ; NAME/KEY:
; LOCATION:
US-08-237-401A-1
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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77.1%; Score 16.2; DB 1; Length 165;
Best Local Similarity 85.7%; Pred. No. 1.1e+02;
Matches 18; Conservative 0; Mismatches 3; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-237-401A-1/C

Sequence 1, Application US/08237401A

Sequence 1, Application US/08237401A

Patent No. 5837441

Patent No. 5837441

Patent No. 5837451

TITLE OF INVENTION: PROTEIN-TYROSINE KINASE GENES

NUMBER OF SEQUENCES: 54

CORRESPONDENCE ADDRESS:

ADDRESSEE: Fish & Richardson P.C.

STREET: 4225 Executive Square, Suite 1400

CITY: La Jolla

STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CONTUTER: IBM PC compatible CONTUTER: IBM PC compatible CONTUTER: IBM PC compatible CONTUTER: IBM PC compatible COSFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/237,401A FILING DATE: 02-MAY-1994
CLASSIFICATION NUMBER: US/08/237,401A FILING DATE: 15-MAY-1994
APPLICATION NUMBER: US 07/884,486
FILING DATE: 15-MAY-1992
APPLICATION NUMBER: US 07/884,486
FILING DATE: 15-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: Halle Ph.D., Lisa A. REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 30,251/007001
TELEPHONE: (619) 678-509
INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
LENGTH: 165 base pairs
TYPE: MOULE: CALABORNES: SINGLE
TYPE: MOULE: CALABORNES: SINGLE
                                                                                07251/007002
ATTORNEY/AGENT INFORMATION:
NAME: Wetherell Ph.D., John R.
REGISTRATION NUMBER: 31,678
REFERENCE/DOCKET NUMBER: 07251/00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 678-5070
TELEFAX: 165 base pairs
LENGTH: 165 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 CCGGGACATCCTCATCCACCC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           76 ccesercárcercaagcacec se
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                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: DNA
IMMEDIATE SOURCE:
CLONE: TYPO-1
                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: CDS
; LOCATION: 1..165
US-08-456-647B-1
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CLONE: Tyro-1
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Ouery Match
77.1%; Score 16.2; DB 2; Length 165;
Best Local Similarity 85.7%; Pred. No. 1.18+02;
Matches 18; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Henderson, Robert A.
APPLICANT: MCNeill, Particle D.
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Hosken, Nancy A.
APPLICANT: Ralos, Michael D.
APPLICANT: Ralos, Michael D.
APPLICANT: Fanger, Gary B.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
TITLE PREPERNICE: 210121.455C6
CURRENT PAPLICATION NUMBER: US/09/480,884A
CURRENT FILING DATE: 2001-08-27
NUMBER OF SEQ ID NOS: 330
SOFTWARE: FREISEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 210121.455C11
CURRENT APPLICATION NUMBER: US/09/643,597
CURRENT FILING DATE: 2000-08-21
NUMBER OF SEQ ID NOS: 369
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 303
LENGTH: 361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-480-884A-303
Sequence 303, Application US/09480884A
Fatent No. 6482597
GENERAL INFORMATION:
APPLICANT: Wang, Tongtong
APPLICANT: Fan, Liqun
                                                                                                                                                                                                                                                                                      Sequence 303, Application US/09643597
Patent No. 6426072
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; CTHER INFORMATION: n = A,T,C or G
US-09-643-597-303
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bangur, Chaitanya S.
Hosken, Nancy
Fanger, Gary R.
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Skeiky, Yasir A.W.
                                                                                                                                                                                                                                                                                                                                                                    Wang, Tongtong
Fan, Liqun
Kalos, Michael D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Li, Samuel X.
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NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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77.1%; Score 16.2; DB 4; Length 361;
Best Local Similarity 85.7%; Pred, No. 1.2e+02;
Matches 18; Conservative 0; Mismatches 3; Indels 0
                                                                                                                                                                                                                                                        Query Match 77.1%; Score 16.2; DB 4; Length 361; Best Local Similarity 85.7%; Pred. No. 1.2e+02; Matches 18; Conservative 0; Mismatches 3; Indels (
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.455C9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Wang, Tongtong
APPLICANT: Ralos, Michael D.
APPLICANT: Ralos, Michael D.
APPLICANT: Alos, Michael D.
APPLICANT: Benger, Nancy A.
APPLICANT: Benger, Gary R.
APPLICANT: Panger, Gary R.
APPLICANT: Panger, Gary R.
APPLICANT: Panger, Gary R.
APPLICANT: NUMBYRION: COMPOUNDS AND METHODS FOR THERAPY
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REPERENCE: 210121.455C8
CURRARN FILING DATE: 2000-04-14
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 303, Application US/09606421B Patent No. 6531315
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; LOCATION: (1) ___(361)
; OTHER INFORMATION: n = A,T,C or G
US-09-542-615A-303
                                                                                                       FEATURE:

NAME/KEY: misc feature

LOCATION: (1)...(361)

CTHER INFORMATION: n = A,T,C or G
US-09-480-884A-303
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APPLICANT: Wang, Tongtong
APPLICANT: Ranos, Michael D.
APPLICANT: Bangur, Chaitanya S.
APPLICANT: Bangur, Chaitanya S.
APPLICANT: Fanger, Gary R.
APPLICANT: Li, Samuel X.
APPLICANT: Li, Samuel X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature
SEQ ID NO 303
LENGTH: 361
TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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US-09-606-421B-303
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LENGTH: 361
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/09/606,421B
CURRENT FILING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 358
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 303
LENGTH: 361
TYPE: DA
TYPE: DB
TYPE:
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Sequence:

Searched:

Sequence Sequence Sequence

Sequence

Sequence

Sequence

Sequence

Sequence Sequence Sequence Sequence Sequence Sequence

Sequence Sequence Sequence

Sequence Sequence Sequence

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Sequence 977, Application US/09016434
| Patent No. 6500938
| GENERAL INFORMATION:
| APPLICANT: Jafice Au-Young
| APPLICANT: Jafice Au-Young
| APPLICANT: Jafice Au-Young
| TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING
| TITLE OF INVENTION: PATHWAY GENE EXPRESSION
| NUMBER OF SEQUENCES: 149
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
| STREET: 3174 POPTER DRIVE
| CITY: PALO ALTO
| STREET: ALLO ALTO
| STREET: ALLO ALTO
| STREET: ALLO ALTO
| STREET: ALLO ALTO
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ZIP: 94304

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS
GURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/016,434
FILING DATE: HEREWITH
CLASSIFICATION NUMBER:
PRICR APPLICATION NUMBER:
FILING DATE: PRICR APPLICATION NUMBER:
FILING DATE: PRICR APPLICATION NUMBER:
                                                                                                                                                                     US-08-706-706-17
US-09-238-471-17
US-09-702-705-95
US-09-736-457-95
US-08-484-661A-38
                                                                                                                 US-09-620-312D-226
                                                                                                                                                                                                                                                                 US-08-656-664-38
PCT-US96-09641-38
US-08-484-661A-36
US-08-656-664-36
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US-08-484-661A-18
US-08-484-661A-22
US-08-484-661A-25
                                                                                                                                                                                                                                                                                                                                                                                                   PCT-US96-09641-10
US-09-257-584-6
US-08-484-661A-7
                                                                                                                                                                                                                                                                                                                                             PCT-US96-09641-36
US-08-484-661A-10
US-08-656-664-10
                                                           US-09-146-053-4
US-07-741-453A-57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-484-661A-28
US-08-484-661A-32
                                                                                                                                                       US-08-706-702-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REPERENCE/DOCKET NUMBER: PA-0002 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                       USA
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US-09-016-434-927
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Command line parameters:
-MODEL=frame+ p2n.model.-DEV=x1h
-Q=/Cgn2_1/USPTO_epool/US08978217/runat_10022004_133826_20481/app_query.fasta_1.1500
-Q=/Cgn2_1/USPTO_epool/US08978217/runat_10022004_133826_20481/app_query.fasta_1.1500
-Q=/Cgn2_1/USPTO_epool/US08978217/runat_10022004_133826_20481/app_query.fasta_1.1500
-LOOPEXTO_-UNITS=bits -STAT=1.FND=1.-MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINIENEO -ALIGN=15
-USDE-CAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINIENEO -MAXENE=200000000
-USDE-CAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINIENEO -LOCGEO -ICPU=3
-NO MMAP -LARGEQUERY -NGG_SCORES= -MAIT -DSPELOCK=100 -LONGLOG
-DEV TIMBOUT=120 -WARN TIMBOUT=30 -THREADS=1 -XGAPDP=10 -XGAPDFXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELDEXT=7
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Sequence 25, Appl
                                                                                                           February 12, 2004, 19:34:21; Search time 4.03329 Seconds (without alignments) 1750.959 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred, No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 27,
Sequence 1, 2
Sequence 1, 2
Sequence 6, 2
Sequence 4, 2
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1: /cgn2_6/prodateal/ina/5A_COMB.seq:*
2: /cgn2_6/prodateal/ina/5B_COMB.seq:*
3: /cgn2_6/prodateal/ina/6A_COMB.seq:*
4: /cgn2_6/prodateal/ina/6B_COMB.seq:*
5: /cgn2_6/prodateal/ina/PCTUS_COMB.seq:*
6: /cgn2_6/prodateal/ina/PCTUS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd
                                                                         OM protein - nucleic search, using frame_plus_p2n model
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US-09-300-958A-27
US-09-300-958A-27
US-08-746-789A-1
US-09-58-68-69-6
US-09-58-43-7-4
US-07-920-281C-1
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US-09-299-141-5
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US-09-554-337-1
                                                                                                                                                                                                                                                                                                                                                                               569978 seqs, 220691566 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                 Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
                                                                                                                                                                                                                               1 KNSSGWKEEEVLQSRN 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                         US-08-978-217-12
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Match Length DB
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8100
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15538
993
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Jatabase :

Result

, Appl.
...ce 6, Appl.
sequence 17, Appl.
Sequence 15, Apr.
Sequence 19

Sequence

Sequence Sequence

Sequence

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WS-09-300-958A-27

Sequence 27, Application US/09300958A

Sequence 27, Application US/09300958A

Sequence 27, Application US/09300958A

Sequence 27, Application US/09300958A

Sequence 37, Application Wichael

APPLICANT: Welsh, John

APPLICANT: Welsh, John

TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TITLE OF INVENTION: Using Same

TITLE OF INVENTION: Using Same

TITLE OF INVENTION: Using Same

TITLE OF INVENTION: 1999-04-27

CURRENT APPLICATION NUMBER: 60/099,070

PRIOR APPLICATION NUMBER: 60/099,070

PRIOR APPLICATION NUMBER: 60/099,070

PRIOR APPLICATION NUMBER: 60/118,624

PRIOR PILING DATE: 1999-02-04

NUMBER OF SEQ ID NOS: 85

SOSTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; LUCRITON: (96)...(1211)
; OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)
; OTHER INFORMATION: protein.
US-09-570-593-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1161 AAAAACTCAAGCGGCTGGAAGGAGGAAGAAGGTTCTCCAGAGTCGGAAC 1208
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Sequence 4 Application US/09570593

Sequence 4 Application US/09570593

Settle No. 6566063

GENERAL INFORMATION:
APPLICANT: Kaufmann, Joerg

APPLICANT: Kaufmann, Joerg

APPLICANT: Kanfmann, Joerg

TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER

FILE REFERENCE: 2300-1556

CURRENT FILING DATE: 2000-05-12

PRIOR APPLICATION NUMBER: 60/134,112

PRIOR APPLICATION NUMBER: 60/134,112

PRIOR FILING DATE: 1999-05-14

NUMBER OF SEQ ID NOS: 13

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4

FAMORIAND APPLICATION WINDOWS OF 13 APPLICATION NOW 1300-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length:
Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Percent Similarity:
Best Local Similarity:
Query Match:
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Pred. No.:
Score:
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LENGTH: 1907
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Sequence 25, Application US/09300958A

Sequence 25, Application US/09300958A

Sequence 25, Application US/09300958A

Sequence 25, Application

APPLICANT: McClelland, Michael

APPLICANT: Welsh, John

APPLICANT: Welsh, John

APPLICANT: Welsh, John

TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TITLE OF INVENTION: Bane: 1999-04-27

CURRENT APPLICATION NUMBER: 06/083,331

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-04-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1999-02-04

NUMBER OF SEQ ID NOS: 85

SOFTWARE: Patentin Ver: 2.0

SEQUID NO. 25
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Matches:
Conservative:
Mismatches:
Indels:
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-978-217-12 (1-16) x US-09-016-434-927 (1-237)
TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 927:
SEQUENCE CHARACTERISTICS:
LENGTH: 237 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.03e-06
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CLONE: 773734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity:
                                                                                                                                                                                                                                                                                                                                                                      Percent Similarity:
Best Local Similarity:
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; LOCATION: (458)
US-09-300-958A-25
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Best Local Similari
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                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match:
DB:
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Sequence 1, Application US/08746789A
Patent No. 5789200
GENERAL INFORMATION: GARDA MARTIN J. Tymms, Christine DeBouck
APPLICANTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1180 AAAACTCAAGCGGCTGGAAGGAGGAAGGAGGTTCTCAAAAAGAGGAAC 1227
                                                                                                                                                                                                   1161 AAAAACTCAAGCGGCTGGAAGGAGGAAGGTTCTCCCAGAGTCGGAAC 1208
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                                                                                                                                                          1 LysAsnSerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative:
Mismatches:
  Conservative:
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                   Mismatches:
Indels:
                                                                                                               US-08-978-217-12 (1-16) x US-09-570-593-4 (1-1907)
                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKilne Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length:
Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HILLING THE TORNET INFORMATION:
NAME: William T. Han
NAME: William T. Han
NEGISTRATION NUMBER: 34,344
REFISENCE TOCKET NUMBER: 34,344
REFISENCE FOOKET NUMBER: 4TG 5
TELECOMMUNICATION INFORMATION: 610 270 4026
TELEFAX: 610 270 4026
INFORMATION FOR SEQ ID NO: 1:
ENGTH: 1920
LENGTH: 1920
TYPE: Nucleic Acid
STRANDEDNESS: Single
TYPE: Nucleic Acid
STRANDEDNESS: Single
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
NO-08-746-789A-1
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) Sequence 6, Application US/09638649

; Patent No. 6563015

; GENERAL INFORMATION:
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Percent Similarity:
Best Local Similarity:
Query Match:
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Query Match:
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STATE: PA
COUNTRY:
                                                                                                                                                                                                                                                 RESULT 5
US-08-746-789A-1
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APPLICANT: Stern, David M.
APPLICANT: Schmidt, Ann Marie
APPLICANT: Schmidt, Ann Marie
APPLICANT: Yan, Shi Du
TITLE OF INVANTION: TRANSGENIC MICE OVER-EXPRESSING RECEPTOR FOR ADVANCED
TITLE OF INVENTION: GLYCATION ENDPRODUCT (RAGE) AND MUTANT APP IN BRAIN AND
TITLE OF INVENTION: USES THEREOF
FILE REPRENCE: 0575/62175
CURRENT APPLICATION NUMBER: US/09/638,649
CURRENT APPLING DATE: 2000-08-14
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length:
Matches:
Conservative:
Mismatches:
Indels:
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Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                               275 AATTCCAGTGGCTGGAAGGAGGAG 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        , ORGANISM: respiratory syncytial virus US-09-554-337-4
                                                                                                                                                                                                                                                                                                                                                                                      2 AsnSerSerGlyTrpLysGluGlu 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 1, Application US/07920281C; Patent No. 5739026
                                                                                                                                                                                                                                                          39.9
46.00
100.00%
100.00%
54.76%
                                                                                                                                                 Patentin Ver. 2.1
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85.71$
50.00$
54.76$
                                                                                                                                                                                                                                                                             Percent Similarity:
Best Local Similarity:
Query Match:
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Best Local Similarity:
                                                                                                                                                                                        TYPE: DNA
ORGANISM: Murine
                                                                                                                                             SOFTWARE: Pate
SEQ ID NO 6
LENGTH: 1348
                                                                                                                                                                                                                                                Alignment Scores:
Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alignment Scores:
Pred. No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-554-337-4
                                                                                                                                                                                                                    JS-09-638-649-6
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us-08-978-217-12.rni

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US-08-466-277-1
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/note= "Semliki Forest Virus complete nucleotide
sequence, presented as a cloned DNA sequence; see
Figure 5."
GENERAL INFORMATION:

APPLICANT: Garcff, Henrik
APPLICANT: Garcff, Henrik
APPLICANT: Liljestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based on
TITLE OF INVENTION: Alphaviruses
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
CITY: Palls Church
STREET: P.O. Box 747
CITY: Falls Church
STREE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   675 GCCACAAACTGGGCCGACGAGGAGGTGTTACAGGCCAGGAAC 716
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/920,281C
FILING DATE: 13-40G-1992
CLASSIFICATION: 435
ATTONNEY/AGENT INPORMATION:
NAME: MUTPHY JT., Gerald M.
REGISTRATION NUMBER: 28,977
REFRENCE/DOCKET NUMBER: 828-103P
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORTATION:
TELECOMMUNICATION INPORMATION:
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TELECOMMUNICATION INPORTATION INPORTATION
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OTHER INFORMATION: /product= "SFV polyprotein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 87..7379
OTHER INFORMATION: /product= "SFV polyprotein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative:
Mismatches:
Indels:
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Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Semliki Forest Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 703-2-1
TELEX: 248345
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 11517 base pairs
"VPE: nucleic acid
"VPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: RNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           46.00
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54.76%
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7421..11179
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LOCATION: 1..11517
OTHER INFORMATION: /
OTHER INFORMATION: /
OTHER INFORMATION: /
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Best Local Similarity:
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LOCATION: 87.
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RESULT 9

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NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STRATE: Wighina
CONDUTER: Wighina
CONDUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: BAP PC COMPATION FOR SYSTEM: US/08/466,277
FILING DATE: 06-Jun-1995
CLASSIFICATION CHIKNOWN>
PRIOR APPLICATION CURKNOWN>
PRIOR APPLICATION CHIKNOWN>
PRIOR APPLICATION CHIKNOWN>
REGISTRATION OF STEMENCE OCKET WINBER: 28,977
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET WUMBER: 28,977
TELEPHONE: 703-241-1300
TELEPRAX: 703-241-1304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 1..11517
OTHER INFORMATION: /label= genome
force= "Semliki Forest Virus complete nucleotide
sequence, presented as a cloned DNA sequence; see
Figure 5."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE DESCRIPTION: /product= "SFV polyprotein" US-08-466-277-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: 87..7379
OTHER INFORMATION: /product= "SFV polyprotein"
Sequence 1, Application US/08466277
Patent No. 6190666
GENERAL INFORMATION:
APPLICANT: Garoff, Henrik
Illjestrom, Peter
TITLE OF INVENTION: DAB Expression Systems Based on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID No: 1:
SEQUENCE CHARACTERISTICS:
TYPE: nucleic acid
STRANDEDRESS: single
TOPOLOGY: linear
MOLECULE TYPE: RNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: CDS
LOCATION: 7421..11179
OTHER INFORMATION: /pr
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85.71$
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ORIGINAL SOURCE:
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Best Local Similarity:
Query Match:
DB:
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Query Match:
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Sequence 1, Application US/09554337

Patent No. 6475780

GENERAL INFORMATION:
APPLICANT: Parrington, Mark
APPLICANT: Li, Xiaomao

ITILE OF INVENTION: ALPHAVIRUS VECTORS FOR PARAMYXOVIRUS VACCINES
CURRENT APPLICATION NUMBER: US/09/554,337

CURRENT FILING DATE: 1030-10-20

PRIOR APPLICATION NUMBER: PC/065,791

PRIOR APPLICATION NUMBER: PCI/CA98/01064

PRIOR FILING DATE: 1997-11-14

PRIOR FILING DATE: 1998-11-13

NUMBER OF SEQ ID NOS: 7

SEQ ID NO 1

SEQ ID NO 1
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                                                                                                                                       675 GCCACAAACTGGGCCGACGAGCAGGTGTTACAGGCCAGGAAC 716
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                                                               3 SerSerGlyTrpLysGluGluGluGluValLeuGlnSerArgAsn 16
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Patent No. 6545200
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Cahoon, Rebeca E.
APPLICANT: Cahoon, Rebeca E.
APPLICANT: Rafalski, J. Antoni
APPLICANT: Rafalski, J. Antoni
TITLE OF INVENTION: STEROL BIOSYNTHETIC ENZYMES
FILE REFERENCE: 1999-12-15
CURRENT FILING DATE: 1999-12-15
EARLIER APPLICATION NUMBER: 60/112,555
EARLIER PLING DATE: 1998-12-16
NUMBER OF SEQ ID NOS: 44
NUMBER OF SEQ ID NOS: 44
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Mismatches:
Indels:
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Mismatches:
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US~08-978-217-12 (1-16) x US-08-466-277-1 (1-11517)
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ) ORGANISM: respiratory syncytial virus US-09-554-337-1
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45.00
78.57%
64.29%
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85.71%
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ORGANISM: Glycine max
US-09-464-535-25
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Best Local Similarity:
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Best Local Similarit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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RESULT 13
US-08-106-925A-12/C
Sequence 12, Application US/08060925A
Fatent No. 543924
Fatent No. 54392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence:p43C-AT-INUS-09-299-141-5
                                                                                                                                                                                                                                                                                                               Sequence 5, Application US/0929141

APPLICANT: FUCTE, TERENCE R.

APPLICANT: SONG, SITONG

APPLICANT: BTRNE, BARRY J.

TITLE OF INVENTION: MATERIALS AND METHODS FOR GENE THERAPY
FILE REFERENCE: 4300.011800

CURRENT APPLICATION NUMBER: US/09/299.141

CURRENT FILING DATE: 1999-04-23

EARLIER PILING DATE: 1998-04-24

NUMBER OF SEQ ID NOS: 13

SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                      661 rccaccaccearcearcaaccarcaaccararracaarccacc 620
                                                                                                2 AsnSerSerGlyTrpLysGluGluGluValleuGlnSerArg 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: KNOBBE, MARTENS, OLSON AND BEAR
STREET: 620 NEWPORT CENTER DRIVE SIXTEENTH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-978-217-12 (1-16) x US-09-299-141-5 (1-7492)
US-08-978-217-12 (1-16) x US-09-464-535-25 (1-993)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
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47.62%
53.57%
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 5
LENGTH: 7492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
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; LOCATION: (180)..(1742) US-09-668-680-8

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6394 AACAGCTCAGGCTGGTTGAGCAACCTTACCTTTAAAGAAGATGTAATTCACCAGAGCAAA 6335
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APPLICANT: Tang, Y. Tom
APPLICANT: Tang, Y. Tom
APPLICANT: Tang, Y. Tom
APPLICANT: Abou, Ping
APPLICANT: Aboundi, Vinod
APPLICANT: Aboundi, Vinod
APPLICANT: Aboundi, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Wu, Andong Jun
APPLICANT: Xu, Chonglun
APPLICANT: Drmanac, Radoje T.
TILLE OF INVENTION: Polypeptides
TITLE OF INVENTION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PL. T. Genes Version 2.0
SEQ ID NO 8
LENGTH: 2763
                SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,925A
FILING DATE: 06-MAY 1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: FUller, Michael L.
REGISTRATION NUMBER: 36,516
REGISTRATION FORSES 1D NO: 12: 12 ELEPAK: 619-235-0376
INFORMATION FOR SEQ ID NO: 12: 5EQUENCE CHARACTERISTICS: 169-235-0376
INFORMATION FOR SEQ ID NO: 12: 5EQUENCE CHARACTERISTICS: 15 ENGTH: 10627 base pairs
TTENDENDESS: single
STRANDENBSS: single
TOTOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-978-217-12 (1-16) x US-08-060-925A-12 (1-10627)
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Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels:
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3-09-668-680-8/c
Sequence 8, Application US/09668680
Patent No. 6436703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-060-925A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           · 6334 AAC 6332
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NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alignment Scores:
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2 - AsnSerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
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                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Asundi, Vinod
APPLICANT: Kue, Aidong J.
APPLICANT: Xue, Aidong J.
APPLICANT: Xu, Chongjun
APPLICANT: Xu, Chongjun
APPLICANT: Xu, Chongjun
APPLICANT: Va, Chongjun
APPLICANT: Va, Chongjun
APPLICANT: Na, Chongjun
APPLICANT: No. 6436703e1 Nucleic Acids and
ITILE OF INVENTION: Polypebtides
FILE REFERENCE: 790CTP2A
CURRENT APPLICATION NUMBER: US/09/668,680
CURRENT APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
FRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 13
SCOFTWARE: DL_FL_genes Version 2.0
             Length:
Matches:
Conservative:
Mismatches:
Indels:
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Search completed: February 12, 2004, 21:50:03 Job time : 9.03329 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps:
                                                                                                                                                                                                                                                                                          Sequence 6, Application US/09668680; Patent No. 6436703
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44.00
66.67%
53.33%
52.38%
                               53.33%
52.38%
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Liu, Chenghua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: CDS
LOCATION: (27)..(2375)
                                     Percent Similarity:
Best Local Similarity: {
Query Match:
DB:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                              0/9-089-899-60-SD
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Pred. No.:
Alignment Scores:
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Sequence 7, Appli Sequence 1, Appli Sequence 37, Appl Sequence 37, Appl Sequence 7864, Ap Sequence 7380, Ap Sequence 7245, Ap

Scoring table: Sequence:

Run on:

1, Appli 1, Appli 9, Appli

Sequence 3 Sequence

sequence 20, Sequence 20, equence 20

Sequence 516, App Sequence 10, Appl

Sequence 13, Appli Sequence 1, Appli

App1i

Sequence 9,

Sequence 7550, Ap Sequence 7788, Ap Sequence 1213, A Sequence 11100, A Sequence 1100, A Sequence 10964, A

Sequence 78, Appl Sequence 1, Appli Sequence 1, Appli

Sequence 3 Sequence 7 Sequence 1 Sequence 3

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RESULT 1
US-09-300-958A-27

i Sequence 27, Application US/09300958A

i Patent No. 6495319

i GENERAL INFORMATION:

APPLICANT: Mcclelland, Michael

APPLICANT: Welsh, John

APPLICANT: Trenkle, Thomas

TILLS OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TILLS OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TILLS OF INVENTION: USing Same

TILLS OF INVENTION: USing Same

TILLS OF INVENTION UNMBER: 60/083,331

PRIOR FILING DATE: 1999-04-27

PRIOR FILING DATE: 1998-04-27

PRIOR APPLICATION NUMBER: 60/098,070

PRIOR APPLICATION NUMBER: 60/098,070

PRIOR APPLICATION NUMBER: 60/118,624

PRIOR FILING DATE: 1999-02-04

PRIOR FILING DATE: 1999-02-04
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US-09-106-6918-23
US-09-106-6918-23
US-09-106-6918-23
US-09-106-6918-23
US-09-106-10-10
US-09-106-10-10
US-09-252-991A-7550
US-09-252-991A-7550
US-09-252-991A-12113
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US-08-252-991A-7865
US-09-252-991A-7865
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US-09-252-991A-7866
US-09-252-991A-7865
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Mismatches:
Indels:
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Matches:
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SOFTWARE: Patentin Ver. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Homo sapiens
US-09-300-958A-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alignment Scores:
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LENGTH: 1907
                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match:
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MODEL=frame+p2n.model -DEV=xlh

MODEL=frame+p2n.model -DEV=xlh

-0=/cgn2 1/USR970 spool/USR9970217/runat 10022004 133826 20481/app query.fasta_1.1500

-DB=18sued_Patents_NA -OFWT=fastap -SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0

-LOOPEXT=0 -UNITS=bits -STRAR=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi

-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15

-NODEE-LOCAL -OUTPWT=pto -NORMeaxt -HEAPSIZE=500 -MINLENS 0 -MAXLENS=200000000

-USER=GOORD -OUTPWT=pto -NORMEAT - 10022004 133926 20481 -NCPU=6 -ICPU=3

-NO MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPELOCK=100 -LONGLOG

-DEV_TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6

-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 27, Appl Sequence 4, Appli Sequence 13.29, A Sequence 13819, A Sequence 139, App Sequence 139, App Sequence 92, Appl Sequence 92, Appl Sequence 92, Appl Sequence 107, Appl Sequence 107, Appl
                                                                                                                                              February 12, 2004, 19:34:21; Search time 21.1748 Seconds (without alignments) 1750.959 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Description
                                                                                                                                                                                                                                                                                                1 NCALEELRLVFGPLGDQLHA......BLLDDGQQASPYHPGSCGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lssued_Patents_NA:*
11. /Ggn2_6/prodateal/ina/5A_COMB.seq:*
12. /Ggn2_6/prodateal/ina/5B_COMB.seq:*
33. /Ggn2_6/prodateal/ina/6B_COMB.seq:*
44. /Ggn2_6/prodateal/ina/6B_COMB.seq:*
55. /Ggn2_6/prodateal/ina/PCTUS_COMB.seq:*
6. /Ggn2_6/prodateal/ina/PCTUS_COMB.seq:*
6. /Ggn2_6/prodateal/ina/PCTUS_COMB.seq:*
  GenCore version 5.1.6
(c) 1993 - 2004 Compugen Ltd.
                                                                                            OM protein - nucleic search, using frame_plus_p2n model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-300-958A-27
US-09-570-593-4
US-09-570-593-4
US-09-252-911A-13819
US-09-252-911A-13819
US-09-344-879A-139
US-09-348-973-139
US-09-215-681-139
US-09-215-681-139
US-09-215-681-139
US-09-215-681-139
US-09-215-681-92
US-09-215-681-92
US-09-215-681-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   569978 seqs, 220691566 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUMMARIES
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                       Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                   US-08-978-217-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query
Match Length DB
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Perfect score:
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4445
73.5
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Result No.

450700

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645 rdrececkéck 656
             81 CysGlyAlaGly 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: Linear ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
Best Local Similarity:
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                                                                             RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      405 AATIGIGGCCTIGAGGAGCIGGGTCTGGTCTTIGGGCCTCTGGGGGACCAACTCCATGCC 464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   41 GluLysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySer 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHisProGlySer 80
                                                                                                                                                                                            21 GlnLeuArgAspleuThrSerSerSerSerSerSerAspGluLeuSerTrpIleIleGluLeuLeu 40
GlnLeuArgAspleuThrSerSerSerSerAspGluLeuSerTrplleIleGluLeuLeu 40
                                                                                                                              41 GlulysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySer 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeuHisAla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (96)...(1211)
OTHER INFORMATION: Human epithelial-restricted with serine box (BSX)
OTHER INFORMATION: protein.
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: KIN, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Harrowe, Greg
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
TITLE OF INVENTION: CANCER
FILE REPRESENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
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Mismatches:
Indels:
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Matches:
                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/09570593
Patent No. 6566063
GENERAL INFORMATION:
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Best Local Similarity:
Query Match:
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LENGTH: 1907
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544 GAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCGGGGAGC
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1 AsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeuHisAla
                                                                                                                                                      US-08-746-789A-1

Sequence 1, Application US/08746789A

Sequence 1, Seplect

Patent No. 5789200

GENERAL INFORMATION:

APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck

TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3

NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 19406-0939
CONDUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
COMPUTER: IBM 486
COMPUTER: MINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746, 789A
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                SEE: SmithKline Beecham Corporation ': 709 Swedeland Road, P.O. Box 1539 King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/746,789A
FILING DATE: No. 5789200ember 15, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: William T. Han
REGISTRATION NUMBER: 34,344
REFERENCE/DOCKET NUMBER: ATG 50024
TELEPHONE: 610 270 5219
TELEPHONE: 610 270 4026
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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NAME/KEY: misc feature
LOCATION: (1)...(521)
OTHER INFORMATION: n = A,T,C or
ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-13819
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16.18%
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73.50
47.95%
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ORGANISM: Homo sapien
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Best Local Similarity:
                                                                                                                                          Similarity:
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                                                                                                                     Percent Similarity:
Best Local Similari
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PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR SEQUENCE 1998-07-27
NUMBER OF SEQUENCE 1998-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AREQUINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: AREQUINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF THE REPRENCE: 107196.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR PILING DATE: 1998-02-18
PRIOR PILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 13819
LENGTH: 801
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Mismatches:
Indels:
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Matches:
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Query Match:
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273 ACTCTCATCAGCAGCTTTTTCAGC-----TTCTTCCAGCTTTTGCAGGGC 229
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APPLICANT: Mitchan, Jennifer L.
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Algate, Paul A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: LAGGOOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C2
CURRENT APPLICATION NUMBER: US/09/404,879A
CURRENT APPLICATION NUMBER: 1999-09-24
NUMBER OF SEQ ID NOS: 393
SOFTWARE: FastSEQ for Windows Version 3.0
SIGO ID NO 139
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Matches:
Conservative:
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                                                             Mismatches:
Indels:
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168 CAAGGAGGCCACCTCAGCCTGTTCCCGGGCCCGCCTTTCTCCCT-CAACTTCTC 110
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273 ACTCTCATCAGCAGCTTTTTCAGC-------TTCTTCCAGCTTTTGCAGGGC 229
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APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
TITLE OF INVENTION: OVARIAN CANCER
FILE REFERENCE: 210121.462C1
CURRENT APPLICATION NUMBER: US/09/338,933
CURRENT PILING DATE: 1999-06-23
NUMBER OF SEQ IO NOS: 312
SOFTWARE: FASTSEQ for Windows Version 3.0
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Matches:
Conservative:
Mismatches:
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; Sequence 139, Application US/09338933
; Patent No. 6488931
; GENERAL INFORMATION;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc feature
LOCATION: (1)...(521)
OTHER INFORMATION: n = A,T,C or G
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ORGANISM: Homo sapien
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US-09-338-933-139
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LENGTH: 521
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168 CAAGGAGGCCACCTCAGCCTGTTCCCGGGCCCGCCTTTCTCCCT-CAACTTCTC 110
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APPLICANT: King, Gordon E.
APPLICANT: King, Gordon E.
APPLICANT: Algare, Paul A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REPERENCE: 210121.465C2
CURRENT APPLICATION NUMBER: US/09/404,879A
CURRENT FILING DATE: 1999-09-24
NUMBER OF SEQ ID NOS: 393
                                                                                               APPLICANT: Mitchan, Jennifer L.
APPLICANT: Frudakis, Tony N.
APPLICANT: Frudakis, Tony N.
APPLICANT: Frudakis, Tony N.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS
TITLE OF INVENTION: OF OVARIAN CANCER
FILE REFERENCE: 210121.463
CURRENT APPLICATION NUMBER: US/09/215,681A
CURRENT PILLEG DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 310
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 139
LENGTH: 521
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Mismatches:
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Matches:
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                                        Sequence 139, Application US/09215681A Patent No. 6528253 GENERAL INFORMATION:
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US-09-404-879A-92/c
US-09-404-879A-92/c
; Sequence 92, Application US/09404879A,
j Eatent No. 6468546
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  , LOCATION: (1)...(521)
; OTHER INFORMATION: n = A,T,C or G
US-09-215-681-139
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ORGANISM: Homo sapien
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Best Local Similarity:
RESULT 8
US-09-215-681-139/c
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354 GAGTICCAICTITICITCAICTITIAAGGCCCGGITITCAAIAACCITCATACCICCTCT 295
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Matches:
Conservative:
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US-09-215-681-92
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US-09-215-681-92/c
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LENGTH: 551
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394 ACTCTCATCAGCAGCTTTTTCAGC------TTCTTCCAGCTTTTTGCAGGC 250
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APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
TITLE OF INVENTION: OVARIAM CANCER
FILE REPERENCE: 210121.462C1
CURRENT APPLICATION NUMBER: US/09/338,933
CURRENT PILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 312
SOFTWARE: PastSEQ for Windows Version 3.0
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Mismatches:
Indels:
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Matches:
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Matches:
      SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 92 LENGTH: 551
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Patent No. 6488931
GENERAL INFORMATION:
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US-09-338-933-92/c
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LENGTH: 551
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354 GAGTICCAICTITICTICAICTITIPAGGCCCGGTITICAATAACCTICATACCTCTCTC 295
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294 ACTCTCATCAGCAGCTTTTTCAGC------TTCTTCCAGCTTTTTGCAGGGC 250
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294 ACTCTCATCAGCAGCTTTTTCAGC------TTCTTCCAGGTTTTGCAGGGC 250
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23 ArgAspLeuThrSerSerSerSerAspGluLeuSerTrpIleIleGluLeuGluLys 42
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                                                                                                                         Sequence 92, Application US/09215681A

Patent No. 652823

GENERAL INFORMATION:

APPLICANT: Mitcham, Jennifer L.

APPLICANT: King, Gordon B.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS

TITLE OF INVENTION: OF COMPOSITIONS AND METHODS FOR DIAGNOSIS

CURRENT APPLICATION NUMBER: US/09/215,681A

CURRENT FILING DATE: 1999-12-17

NUMBER OF SEQ ID NOS: 310

SOFTWARE: FASTSEQ for Windows Version 3.0
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SEQ ID NO 107
LENGTH: 555
 SEQ ID NO 107
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Sequence 107, Application US/09404879A

Parent No. 6468346

GENERAL INFORMATION:
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Algate, Paul A.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.462C2

CURRENT APPLICATION NUMBER: US/09/404,879A

CURRENT PILING DATE: 1999-09-24

NUMBER OF SEQ ID NOS: 33.0
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APPLICANT: Mitchem, Jennifer Lynn
APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
TITLE OF INVENTION: OVARIAN CANCER
TITLE OF INVENTION: OVARIAN CANCER
CURRENT APPLICATION UNDER: US/09/338,933
CURRENT PILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 312
SOFTWARE: FASELSEQ FOR Windows Version 3.0
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Mismatches:
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Patent No. 6488931
GENERAL INFORMATION:
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81 CysGlyAla 83
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US-09-404-879A-107
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Patent No. 6528253
GENERAL INPORMATION:
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Frudakis, Tony N.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS
TITLE OF INVENTION: OF OVARIAN CANCER
FILE REFERENCE: 210121.463
CURRENT APPLICATION NUMBER: US/09/215,681A
CURRENT FILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 310
SOFTWARE: FREUSEQ for Windows Version 3.0
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Mismatches:
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Conservative:
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72.00
39.81%
30.10%
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72.00
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16.18%
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LENGTH: 555
TYPE: DNA
CRGANISM: Homo sapien
US-09-338-933-107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapien
                                                                                                                                                   Percent Similarity:
Best Local Similarity:
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US-09-215-681-107/c
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Search completed: February 12, 2004, 21:49:58 Job time : 24.1748 secs
Gaps:
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                                                                     298 ACTCTCATCAGCAGCTTTTTCAGC------TTCTTCCAGCCTTTTGCAGGGC 254
358 GAGTICCATCITICTICATCITITAAGGCCCGGITITICAATAACCITCATACCICICIC 299
                                                                                                                                                253 AGTGGCCAGGCGCTCCTGAGCACGGTCCAGCTCTTCAACCAGCTGGATCCTACGTT 194
                                                                                                                                                                                   ---ProPheAspGlnGlySerProPheAlaGlnGluLeuLeu 67
                                                                                                                                                                                                                                                              68 AspAspGlyGlnGlnAlaSerProTyrHis---------ProGlySer 80
                                     23 ArgAspLeuThrSerSerSerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLys 42
                                                                                                              43 AspGlyMetAla------54
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23, Application US/08306691B
Patent No. 5734039
GENERAL INFORMATION:
APPLICANT: Calabretta, Bruno
APPLICANT: Skorski, Tomasz
TITLE OF INVENTION: ANTISENSE
TITLE OF INVENTION: OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
NUMBER OF SEQUENCES: 55
CORRESPONDENCE ADDRESS:
ADDRESSE: Seidel, Gonda, Lavorgna & Monaco, P.C.
STREET: Two Penn Center, Suite 1800
CITY: Philadelphia
STATE: Philadelphia
STATE: Philadelphia
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Mismatches:
Indels:
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MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
COMPUTER: IBM PS/2
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/306,691B
FILING DATE: September 15, 1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Monaco, Daniel A.
REGISTRATION UNIBER: 30,480
REFERENCE/DOCKET NUMBER: 8321-8
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-5383
TELERA: (215) 568-549
TELERA: No. 5734039e
INFORMATION FOR EQQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 2301 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 2301 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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COUNTRY: U.S.A.
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Best Local Similarity:
Query Match:
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US-08-306-691B-23
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388 CAAGGAGGCCACCTCAGCCTGTTGCCGGGCCGGCCTTTCTCCCT-CAACTTCTC 330
                                      329 gcragaAggcgcrcaccrccrcchcracArcarcrgccrgcrgcrgcagaAccrgdArcr 270
                                                                                                                                                     493 ACTCTCATCAGCAGCTTTTTCAGC------TTCTTCCAGCTTTTGCAGGGC 449
                                                                                                                                                                                                                                                       448 AGTGGCCAGGCGCCCCTGAGCACGGTCCAGCTCTTCAACCAGCTGGATCCTACGGTT 389
                                                                                                                                                                                                              43 AspGlyMetAla-------54
                                                                                                                                                                                                                                                                                                                                                                                        80
                                                                                                                         23 ArgAspLeuThrSerSerSerSerAspGluLeuSerTrplleIleGluLeuLeuGluLys 42
                                                                                                                                                                                                                                                                                                     -----ProGlySer
                                                                                                                                                                                                                                                                                                                                                                                            68 AspAspGlyGlnGlnAlaSerProTyrHis-------
US-08-978-217-7 (1-84) x US-08-306-691B-23 (1-2301)
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Sequence 1, Appli
Sequence 3437, Ap
Sequence 12009, A
Sequence 8007, Ap
Sequence 8007, Ap
Sequence 1, Appli
Sequence 1, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 9092, Ap
Sequence 9092, Ap
Sequence 9082, Ap
Sequence 8997, Appli
Sequence 8997, Appli
Sequence 1880, Ap
Sequence 1881, Ap
Sequence 1881, Ap
Sequence 1881, Appli
Sequence 1881, Appli
Sequence 1083, Appli
Sequence 1083, Appli
Sequence 1083, Appli
Sequence 11, Appli
                                                                                     February 12, 2004, 21:46:16; Search time 14.4951 Seconds (without alignments) 7673.534 Million cell updates/sec
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                                                                                                                                                                               Description
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-746-789A-1

US-08-313-224A-3437

US-08-242-671-1206

US-09-252-991A-1206

US-09-252-991A-8097

US-09-252-991A-8097

US-09-185-370-1

US-09-185-370-1

US-09-185-370-1

US-09-111-444-4

US-09-252-991A-8781

US-09-252-991A-8995

US-09-252-991A-8995

US-09-252-991A-8995

US-09-252-991A-8995

US-09-252-991A-8995

US-09-252-991A-8995

US-09-252-991A-8995

US-09-252-991A-8995

US-09-252-991A-8995

US-09-252-991A-8995

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US-09-252-991A-8995

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US-09-252-991A-8995
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US-09-252-991A-7749
US-09-252-991A-7739
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-300-958A-27
US-09-570-593-4
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                                                                                                                                                                                                                                                                                                                                                             Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                        OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                            IDENTITY NUC
Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                    length: 0
length: 200000000
                                                                                                                                                US-08-978-217-6
252
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query
Match Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   741
918
1650
2145
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2161
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                                                                                                                                                Title:
Perfect score:
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Maximum DB seq
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                                                                                                                                                                               Sequence:
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ALIGNMENTS

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Sequence 27, Application US/09300958A

Patent No. 649519

GENERAL INFORMATION:
APPLICANT: McClelland, Michael
APPLICANT: McClelland, Michael
APPLICANT: McClelland, Michael
TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of
TITLE OF INVENTION: Using Same
TITLE OF INVENTION: Using Same
FILE REFERENCE: P-PH 3457
CURRENT APPLICATION NUMBER: US/09/300,958A
CURRENT FILING DATE: 1999-04-27
CURRENT PAPLICATION NUMBER: 60/083,331
PRIOR FILING DATE: 1998-04-27
PRIOR FILING DATE: 1998-08-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
US-09-300-958A-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-300-958A-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (96)...(1211)
OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)
OTHER INFORMATION: protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Application US/08746789A
Patent No. 5789200
GENERAL INFORMATION:
APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELP3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 252; DB 4; Length 1907; 100.0%; Pred. No. 5e-59; tive 0; Mismatches 0; Indels 0
            Sequence 4, Application US/09570593
| Patent No. 6566063
| GENERAL INFORMATION:
| APPLICANT: Kalfmann, Joerg
| APPLICANT: Kin, Hong
| APPLICANT: Kin, Hong
| TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN TITLE OF INVENTION: CANCER;
| FILE REFERENCE: 2300-1556
| CURRENT FILING DATE: 2000-05-12
| PRIOR PILING DATE: 2000-05-14
| NUMBER OF SEQ ID NOS: 13
| SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM 486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: SmithKline Beecham Corporation STREET: 709 Swedeland Road, P.O. Box 1539 CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         241 TGTGGCGCAGGA 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches 252; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sest Local Similarity
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COUNTRY:
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US-09-570-593-4
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                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                       PE: DNA
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APPLICANT: Laigudi, Raghunath V.
APPLICANT: Laigudi, Raghunath V.
APPLICANT: Laigudi, Raghunath V.
APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN EAR FILE REFERENCE: PL-0017 US
CURRENT APPLICATION NUMBER: US/09/313,294A
CURRENT PILING DATE: 1999-05-14
NUMBER OF ERQ ID NOS: 7600
SSOFTWARE: PERL Program
SEQ ID NO 3437
LENGTH: 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         484 CAGCTGCGAGACTCCAGCTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG 543
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    544 GAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGC 603
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; OTHER INFORMATION: Incyte ID No. 6476212 700611585H1
US-09-313-294A-3437
OPERATING SYSTEM: WINDOWS FOR WORGROUPS
SOTTWARE: MICROSOTT WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746,789A
FILLING DATE: No. 5789200ember 15, 1996
CLASSIFTCATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER:
FILLING DATE:
APPLICATION NUMBER: 34,344
REGISTRATION NUMBER: 34,344
REGISTRATION NUMBER: 34,344
RETEMENCE/DOCKET NUMBER: ATG 50024
TELEPHONE: 610 270 5219
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 100.0
Matches 252; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single
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STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: No
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ORGANISM: Zea mays
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US-08-746-789A-1
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ORGANISM: Pseudomonas aeruginosa
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US-09-252-991A-12066
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and Uses Thereof in regulating Gene Expression and in the
Treatment of AIDS
                                                                                                                                                           42 GGGGGACCAACTCCATGCCCAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAG 101
                                                                                                                                                                                                                                      60 GGTGGAGCCACTGCTGGCGGAGTCCTCCGACGCTTCTCCCATGTTCCCGATCCGCTTCCC 119
                                                                                                                                                                                                                                                                                                                      102 TIGGATCATIGAGCIGCIGGAGAAGGAIGGCAIGGCCTICCAGGAGGCCCIAGACCCAGG 161
                                                                                                                                                                                                                                                                                                                                                                                             120 GCAGATCTGGGAGTTCTACAAGAAGGCGGTGGTCCTTCTGGACGGCGAGGAGGTTGA 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             162 GCCCTTTGACCAGGCCCCCTTTGCCCAGGAGCTGCTGGACGACGACGGCAAGCCAG 221
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15.4%; Score 38.8; DB 1; Length 5173;
Best Local Similarity 50.5%; Pred. No. 0.16;
Matches 94; Conservative 0; Mismatches 92; Indels 0
    DB 4; Length 276;
Query Match
15.5%; Score 39; DB 4; Length 276
Best Local Similarity 49.3%; Pred. No. 0.07;
Matches 102; Conservative 0; Mismatches 105; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/242,677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        222 CCCCTACCACCCGGCAGCTGTGGCGC 248
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ATTORNEY/ABENT INFORMATION:
NAME: Mayfield, Denise L.
REGISTRATION NUMBER: 33,732
REFERENCE/DOCKET NUMBER: UTSD:401
TELECOMMUNICATION INFORMATION:
TELECHONE: 713-787-14400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433
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IBM PC compatible
YSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-242-677-1/c
; Sequence 1, Application US/08242677
; Patent No. 5677143
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APPLICANT: Gaynor, Richard B
APPLICANT: W., Foon W.
TITLE OF INVENTION: Cellular Nu
TITLE OF INVENTION: and Uses Th
TITLE OF INVENTION: Treatment o
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
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SOFTWARE: Patentl
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LOCATION: 1..4863
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48 CCAACTCCATGCCCAGCTGCGAGACCTCCAGCTCTTCTGATGAGCTCAGTTGGAT 107

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APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1999-02-18
PRIOR PLICATION NUMBER: US 60/074,190
PRIOR APPLICATION NUMBER: US 60/094,190
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 12066
LENGTH: 741
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Sequence 12009, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: NUCLEIC AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT PILING DATE: 1999-02-18
PRIOR PILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-07-19
PRIOR FILING DATE: 1999-07-27
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Best Local Similarity 47.6%; Pred. No. 0.51;
Matches 107; Conservative 0; Mismatches 118; Indels
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Matches 104; Conservative
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US-08-852-743-1
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Sequence 8097, Application US/09252991A
Sequence 8097, Application US/09252991A
Sequence 8097, Application US/09252991A
Sequence 8097, Application US/09252991A
SENERAL INFORMATION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS;
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS;
TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
CURRENT EXPELICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
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                                                                                                                                                              14.4%; Score 36.2; DB 4; Length 918; 47.6%; Pred. No. 0.53;
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                                                                 TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
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NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                              Query Match
Best Local Similarity 47.6
Matches 107; Conservative
                                                                                                              US-09-252-991A-12009
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US-09-252-991A-8097
                    SEQ ID NO 12009
LENGTH: 918
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RESULT 9 US-09-252-991A-8205

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Sequence 8205, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION:
NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
FILE REFERENCE: 107196.136
CURRENT FILING DATE: 1999-02-18
PRIOR PLILING DATE: 1999-02-18
PRIOR PLILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 8205
LENGTH: 2145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1576 CCAGCCCTGGGCGCCGCGTCCACCGGCGATCTCGGCGGCGGCCAGGCACAGGCGGCGAA 1635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1636 GCGCAGGTGGTTGTCCGGCCAGTCGTTGCCCTGGTCGTCCATGTACGGCGTACCCTCGCG 1695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1516 GGCCATGTAGGCCGGCGTCAGCGCCGACGGCCAGTCGTTGGCGTGCACCAGGCCCGGTTG 1575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                67 CGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTGGAGAAG 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            127 GATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGCCCCTTT 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7 GCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCCCAGCTG 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 2145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14.4%; Score 36.2; DB 4; 47.9%; Pred. No. 0.66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Force, Thomas
APPLICANT: Kyriakis, John M.
APPLICANT: Pombo, Cella M.
APPLICANT: Pombo, Cella M.
APPLICANT: Bonyentre, Joseph
TITLE OF INVENTION: SOK-1 AND METHODS OF USE
NUMBER OF SEQUENCES: 10
CORRESOEDE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE PORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: Diskette
COMPUTER: DISKETTE
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COMPUTER: DISKETTE
COMPUTER: DISKETTE
COMPUTER: DATE:
COMPUTER: DISKETTE
COMPUTER: DATE:
APPLICATION NUMBER: US/08/852,743
FILING DATE:
APPLICATION NUMBER: 60/016,774
FILING DATE:
APPLICATION DATA:
APPLICATION NUMBER: 60/016,774
FILING DATE: 7-MAY-1996
ATTORNEY/AGENT INFORMATION:
NAME: FRASET, JANIS K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/08852743
Patent No. 5830699
GENERAL INFORMATION:
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us-08-978-217-6.rni

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RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       175 GAGGAGCTCTTCACCAAGCTCGACCGCATTGGCAAGGGCTCGTTTGGGGAAGGTCTACAAG 234
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                                                                                                                                                                                                                                                                                                                                                        13.9%; Score 35; DB 2; Length 1975; 48.3%; Pred. No. 1.4;
                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 1.4;
0; Mismatches 105; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Force, Thomas
APPLICANT: Kyriakis, John M.
APPLICANT: Pombo, Celia M.
APPLICANT: Bonventre, Joseph
TITLE OF INVENTION: SOK-1 AND METHODS OF USE
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 02110-2804
COMPUTER READABLE PORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COPENATING SYSTEM: Windows95
SOFTWARE: PeacESQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/185,370
FILING DATE:
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00786/327001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-5070
TELEFAX: 200154
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1975 base pairs
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APPLICATION NUMBER: 08/852,743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/09185370 Patent No. 6093560 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                            Coding Sequence 127...1404
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                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 48.3
                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                   MOLECULE TYPE: CDNA
                                                                                                                                                                                                                     linear
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                                                                                                                                                                                                                                                                          NAME/KEY:
LOCATION:
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US-09-185-370-1
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Matches
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46 GACCAACTCCATGCCCAGCTGCGAGACCTCCACTTCCAGCTCTTCTGATGAGCTCAGTTGG 105
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13.9%; Score 35; DB 3; Length 1975; 48.3%; Pred. No. 1.4;
                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches 105; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/08712709
Patent No. 5863780
GENERAL INFORMATION:
APPLICANT: AL-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION NUMBER: US/08/712,709
FILING DATE: Filed Herewith
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0118 US
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TELEPHONE: 415-855-0555
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                               TELEX: 200154
INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 1975 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 2161 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                      Coding Sequence 127...1404
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                                                                                                                                                                                                                                                                                                                                                                                                        98; Conservative
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Best Local Similarity
Matches 98; Conserv
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LOCATION:
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CLONE: Consensus
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Best Local Similarity
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                 US-09-111-444-4
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US-09-541-228-4
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                                                                                                                                                                                                                                                                                                    106 ATCATTGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCC 165
                                                                                                                                                                                                                                                                                                                                                                            346 GGCATCGATAACCACACAAAGGAGGTGGTGGCCATCAAGATCATCGACCTGGAGGAGGCC 405
                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
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1 Similarity 48.3%; Pred. No. 1.4; 
98; Conservative 0; Mismatches 105; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94304

COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COPERATING SYSTEM: DOS
SOFTWARE: FESTERED, Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/111,444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PF-0118 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                      226 TACCACCCGGCAGCTGTGGCGC 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           466 TACATCACCCGCTACTTTGGCTC 488
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APPLICATION NUMBER: 08/712,709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4, Application US/09111444 Patent No. 6045792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELBERAY: 415-845-4166
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 2161 base pairs
TYPE: mucleic acid
STRANDEDNESS: single
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
IMMEDIATE SOURCE:
                                                                                                                                                 Query Match
Best Local Similarity
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                                                                       LIBRARY:
CLONE: Consensus
US-08-712-709-4
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IMMEDIATE SOURCE:
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46 GACCAACTCCATGCCCAGCTGCGAGACCTCCAGCTCCAGCTCTTCTGATGAGCTCAGTTGG 105
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                                                                                                                                                                                                                                                            346 GGCATCGATAACCACACACAGAGGAGGTGGCCATCAAGATCATCGACCTGGAGGAGGCC 405
                                                                                                                                                                                                                                                                                                            406 GAGGÁTGÁGATÓGAGGACATOCAGCAGGAGATCACTGTÓCTÓAGTÓAGTGCGACAGCÓÓC 465
                                                                                                                                                                                                             106 ATCATTGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCC 165
                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 35; DB 3; Length 2161; Pred. No. 1.4;
          13.9%; Score 35; DB 3; Length 2161;
48.3%; Pred. No. 1.4;
ative 0; Mismatches 105; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
INMER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/541,228
                                                                                                                                                                                                                                                                                                                                                                                                             226 TACCACCCCGGCAGCTGTGGCGC 248
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PILING DATE:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION TOWBER: 08/712,709
FILING DATE:
ATTOMEY/ABOUT INFORMATION:
NAME: Bilings, Lucy J
REGISTRATION NUMBER: 36,749
REPERENCE/DOCKET NUMBER: PF-011
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/09541228 Patent No. 6232077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13.9%;
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 2161 base pairs
Query Match
Best Local Similarity 48.39
Matches 98, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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RESULT 15

US-09-222-991A-8781/C

Squence 8781, Application US/09252991A

Patent No. 6551795

GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

TITLE OF INVENTION: ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: 1009-02-18

CURRENT APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 8781

LENGTH: 621
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                                       46 GACCAACTCCATGCCCAGGTGCGAGACCTCCAGCTCTTCTGATGAGCTCAGTTGG 105
                                                                                106 ATCATTGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCC 165
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0; Mismatches 105; Indels
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                                                                                                                                                                                                                                                                                                                     226 TACCACCCCGGCAGCTGTGGCGC 248
                                                                                                                                                                                                                                                                                                                                                                 466 racarcaccecraciriescre 488
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ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-8781
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Best Local Similarity 54.84
Matches 69; Conservative
  Matches 98; Conservative
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Sequence 44, Appl Sequence 44, Appl Sequence 44, Appl Sequence 338, Appl Sequence 10, Appli Sequence 10, Appli Sequence 17, Appli

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RESULT 1

US-09-300-958A-27

i Sequence 27, Application US/09300958A

i Ratent No. 6495319

i GENERAL INFORMATION:

APPLICANT: MccIelland, Michael

APPLICANT: MccIelland, Michael

APPLICANT: MccIelland, Michael

APPLICANT: Mclalland, Michael

APPLICANT: Mclalland, Michael

APPLICANT: Welsh, John

TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

TITLE OF INVENTION: Using Same

FILE REFERENCE: P-PH 3457

CURRENT FILING DATE: 1998-04-27

PRIOR PLICATION NUMBER: 60/093,331

PRIOR PLICATION NUMBER: 60/093,331

PRIOR PLING DATE: 1998-04-27

PRIOR PLING DATE: 1998-04-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

PRIOR FILING DATE: 1998-08-27

SOFTWARE OF SEQ ID NOS: 85

SOFTWARE PATENTION VET: 2.0
                              US-09-439-313-44
US-09-1352-616A-44
US-09-03-1352-616A-44
US-09-039-913-8
US-09-039-913-8
US-09-039-913-6
US-09-039-913-6
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US-09-05-913-6
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-WODEL=frame+_p2n.model -DEV=xlh
-Q=/CQMI_21/VBPTO spool/U308978217/runat_10022004_133826_20481/app_guery.fasta_1.1500
-Q=/CQMI_21/VBPTO spool/U308978217/runat_10022004_133826_20481/app_guery.fasta_1.1500
-DB=ISSUEG_PATEPTO START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MXX=100 -THR_MIN=0 -ALIGN=15
-MODE_LCCAL -OUTFWTT=pto -NOFM=ext -HEAPSIZE=500 -MINLENS -NAXIEN=200000000
-USER=LOCAL -OUTFWTT=pto -NOFM=xt 10222004_133826_20481 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DBV_TIMEOUT=120 -WARN_TIMEOUT=30 -THRADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELDP=6 -DELEXT=7
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Sequence 282, App
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Sequence 2, Appli
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Sequence 44, Appli
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Sequence 4, Appli
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                                                                                                                                                                                                                                                                    February 12, 2004, 19:34:21; Search time 93.522 Seconds (without alignments) 1750.959 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MAATCEISNIFSNYFSAMYS......YKFGKNSSGWKEEEVLQSRN 371
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2: /cgm2_6/ptcdata/1/ina/5B_COMB.seq:*
3: /cgm2_6/ptcdata/1/ina/6A_COMB.seq:*
4: /cgm2_6/ptcdata/1/ina/6B_COMB.seq:*
5: /cgm2_6/ptcdata/1/ina/PCTUS COMB.seq:*
6: /cgm2_6/ptcdata/1/ina/PcTUS COMB.seq:*
                                  GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                      OM protein - nucleic search, using frame_plus_p2n model
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US-09-570-593-4
US-08-746-798A-1
US-09-620-405B-282
US-09-339-882-82
US-09-333-86-282
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US-09-333-86-282
US-09-009-913-3
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Maximum Match 100%
Listing first 45 summaries
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Kgapop 10.0 , Kgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Maximum DB seq length: 200000000
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: LOCATION: (96)...(1211)

: OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)

: OTHER INFORMATION: protein.

US-09-570-593-4
                                                                                                                                                                                    APPLICANT: Kaufmann, Joerg
APPLICANT: Kaufmann, Joerg
APPLICANT: Kan, Hong
APPLICANT: Harrowe, Greg
APPLICANT: Harrowe, Greg
TITLE OF INVENTION: CANCER
FILE REFRESENCE: 2300-1556
CURRENT APPLICATION NUMBER: US/09/570,593
CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR PILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FREUSEQ for Windows Version 4.0
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
1176 TGGAAGGAAGAAGATTCTCCAGAGTCGGAAC 1208
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                                                                RESULT 2
US-09-570-593-4
; Sequence 4, Application US/09570593
; Patent No. 6566063
; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Percent Similarity:
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SEQ ID NO 4
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GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHis 180
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                                                                                                                                                                                                                                                                              241 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys 260
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                                                                                                                                                                                                                             CCCACTGATGGCAAGCTCTTCCCCAGCGATGGTTTTCGTGACTGCAAGAAGAGGGGGATCCC
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                                                                                                                                       ThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySerAspValAspLeuAsp
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                                                                     181 ProGlySerCysGlyAlaGlyAlaProSerProGlySerSerAspValSerThrAlaGly
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Patent No. 5789200
GENERAL INFORMATION:
APPLICANT: Ismail Kola, Martin J. Tymms, Christine DeBouck
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
CITY: King of Prussia
STATE: PA
COUNTRY: USA
ZIP: 19406-0939
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ZIP: 19406-0939

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM 486
COMPUTER: IBM 486
COPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MCROSOFT WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/746,789A
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
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Mismatches:
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Matches:
                       MATE: MILLIAMI 1. MATE:

REGISTRATION NUMBER: 34,344

REFERENCE/DOCKET NUMBER: ATG 50024

TELECOMMUNICATION INFORMATION:

TELEPHONE: 610 270 5219

TELEPHONE: 610 270 4026

INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:

LENGTH: 1920

TYPE: Nucleic Acid

STRANDEDNESS: Single

TOPOLOGY: Linear

GS-08-746-789A-1
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1980.00
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William T. Han
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Query Match:
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us-08-978-217-2.rni

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US-08-978-217-2 (1-371) x US-09-620-405B-282 (1-502)
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                                                                                                                                                                                                                                                                                                                                                                                344 ArgValAspGlyArgArg 349
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US-09-620-405B-282
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US-09-620-405B-282
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LENGTH: 502
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                                                                                                                                                          AspileLeuileHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 300
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                             835 AAGCACGGGAAGCGGGAAACGAGGCCCGGAAAGCTGAGCAAAGAGAAAGAGTACTGGGGACTGT 894
                                                                                280
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APPLICANT: Yuqui, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.47003
CURRENT APPLICATION NUMBER: US/09/389,681A
CURRENT FILING DATE: 1999-09-02
NUMBER OF SEQ ID NOS: 463
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 282
LENGTI: 502
241 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys
                                                                                261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg
                                                                                                                895 CTCGAGGGCAAGAAGAGCAAGCACGCCCCAGAGGCACCCACCTGTGGGAGTTCATCCGG
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Matches:
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US-09-189-681-282
Sequence 282, Application US/09389681A
Patent No. 6518337
GENERAL INFORMATION:
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US-09-389-681-282
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324 MetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGluIleLeuGlu 343
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                                                                                                                                                                                                                                303 ATCCACCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAAGGCGTTTC 362
                                                                                                                                                                                                                                                                                                          304 LysPheLeukrgSerGluklaValAlaGlnLeuTrpGlyGlnLysLysLysBysSerAsn 323
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                                                               283
                                                                                             243 AAGAAGAGGCACGCGCCCAGAGGCACCCACTGTGGGAGTTCATCCGGGACATCCTC 302
                                                                                                                                                                                        284 IleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGluGlyValPhe 303
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                                                               264 LysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheIleArgAspIleLeu
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APPLICANT: Jiang, Yuqiu
APPLICANT: Jiang, Yuqiu
APPLICANT: Jiang, Yuqiu
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Hepler, William T.
TITLE OF INVENTION: DAGNOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REPERRENCE: 210121.47008
CURRENT APPLICATION WUMBER: US/09/620,405B
CURRENT FILING DATE: 2000-07-20
NUMBER OF SEQ ID NOS: 495
SOFTWARE: FRAESEQ for Windows Version 3.0
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Matches:
Conservative:
Mismatches:
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304 LysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysAsnSerAsn 323
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Mismatches:
Indels:
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Matches:
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; Patent No. 6579973
; GENERAL INFORMATION:
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Best Local Similarity:
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SEQ ID NO 282
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264 LysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheIleArgAspIleLeu 283
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APPLICANT: Vidiu, Jiang
APPLICANT: Mitchem, Jennifer L.
APPLICANT: Mitchem, Jennifer L.
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APPLICANT: Mitchem, Jennifer L.
TITLE OF INVENTION: DIAMOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C2
CURRENT APPLICATION NUMBER: US/09/339, 338A
CURRENT APPLICATION NUMBER: US/09/339, 338A
CURRENT FILING DAIE: 1999-06-23
NUMBER OF SEQ ID NOS: 315
SOFTWARE: FRAESEQ for Mindows Version 3.0
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CRGANISM: Homo sapiens
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Query Match:
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303 ATCCACCGGGAGCTCAACGAGGGCCTCATGAAGTGGGAGAATCGGCATGAAGGGCTTTC 362
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APPLICANT: Jiang, Yuqui
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Midangchun
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C4
CURRENT APPLICATION NUMBER: US/09/433,826B
CURRENT PLING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 1999-11-03
SOFTWARE: FastSEQ for Windows Version 3.0
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REGISTRATION NUMBER: 36,677
REFERENCE DOCKET NUMBER: SEQ-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
TELEFAX: 650-327-3231
                                                                                                                                    344 ArgvalAspGlyArgArg 349
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SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US:
FILING DATE: 21-JAN-199
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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423 ATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAGATCCTGGAA 482
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patent No. 6586572

GENERAL INPORMATION:
APPLICANT: Jang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSES OF BREAST CANCER
FILE REFERENCE: 210121.470c7
CURRENT FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 489

SOFTHARE: FRSISEQ for Windows Version 3.0
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Conservative:
Mismatches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-604-287A-282
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LENGTH: 502
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DB:
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Matches:
Conservative:
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Indels:
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                                                                                                                                                              Sequence 2, Application US/09009913
| Batent No. 6087485
| GENERAL INFORMATION:
| TITLE OF INVENTION: Asthma Related Genes
| NUMBER OF SEQUENCES: 339
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Bozicevic & Reed, LLP
| STREET: 285 Hamilton Ave, Suite 200
| CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94301
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPREATING SYSTEM: DOS
SOPTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UMBER: US/09/009,913
21-JAN-1998
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GENERAL INFORMATION:

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255 GGGTTTTTTGGAGGCCAGTGGAATTCATCCTCAGTACTGGACCAAGTACCAGGTG 314
                                                                                             315 TGGGAGTGGCTCCAGCACCTCCTGACAGCTGGATGCCAATGTATCCTTTC 374
                                                                                                                                                                                                                             113 ValPheGlyProLeuGlyAspGlnLeuHisAlaGlnLeuArgAspLeuThr----- 129
                                                                                                                                                                                                                                                   150 PheGlnGluAlaLeuAspProGlyProPheAspGlnGlySerProPheAlaGlnGluLeu 169
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                                                                                                                                                  93 SerArgCysAspMetAspGlyAlaThrLeuCysAsnCysAlaLeuGluGluLeuArgLeu 112
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                                                                       73 LeuAspTrpIleSerTyrGlnValGluLysAsnLysTyrAspAlaSerAlaIleAspPhe 92
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; Sequence 3, Application US/09009913
; Patent No. 6087485
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 59 TrpLeuGlyGluGlnProGlnPheTrpSerLysThrGlnValLeuAspTrpIleSerTyr 78
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Mismatches:
Indels:
                                                                                                                                                                                                                                                              SOFTWARE: FASTESTO FOR Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/nno ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-978-217-2 (1-371) x US-09-009-913-3 (1-5510)
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Matches:
APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bozicevic & Reed, ILP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                          UMBER: US/09/009,913
21-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               36,677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: SE
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 5510 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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555.50
51.24%
39.13%
28.06%
                                                                                                                                                                                                                                                            IBM Compatible
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ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: CDNA
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                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                               USA
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                                                                                    196 ValSerThrAlaGlyThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySer 215
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COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 4, Application US/09009913
Patent No. 6087485
GENERAL INFORMATION:
APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
TUTHE OF INVENTION: Asthma Related Genes
CORRESPONDENCE ADDRESS:
ADDRESSEE: Boatcevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
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21-JAN-1998
                                          635 ACTGAGCCT------
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                                                                                                                                                                                                                                                              236 LysLys------
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MEDIUM TYPE: Diskett
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CLASSIFICATION:
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                  Length:
Matches:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: SEQ
TELECOMMUNICATION:
TELEPHONE: 650-327-3231
                                                                                                                                     INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 5667 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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TELEX:
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Query Match:
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Sequence 44, Application US/09030607; Patent No. 6262245; GENERAL INFORMATION: GAPLICANT: Au Jiangchun APPLICANT: Xu, Jiangchun APPLICANT: Dillon, Davin C. TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO
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                                                                                    91 AspPheSerArgCysAspMetAspGlyAlaThrLeuCysAsnCysAlaLeuGluGluLeu 110
                                                                                                         730 ACCCGGGCGGCAGGGACGGCGGGGCANCTCCTCTACAGCAACTTGCAGCATCTGAAGTGG 671
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Sequence 44, Application US/09020956

Sequence 44, Application US/09020956

Sequence 44, Application US/09020956

GENERAL INFORMATION:
APPLICANT: Dillin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FOR NUMBER OF SEQUENCES: 178

CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                         .041 TTGAACCCAGACAAGAACCCAGGATTAATAAAATGGGAAGACCGATCTGAGGGGCGTCTTC 1100
                                                                                                                            1101 AGGITCITGAAATCAGAGGCAGTGGCTCAGCTATGGGGTAAAAAGAAGAACAACAGCAGC 1160
                                                                                                                                                                                                              1161 ATGACCTATGAAAAGCTCAGCCGAGCTATGAGATATTACTACAAAAGAGAAATACTGGAG 1220
                                                                                  304 LysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLysAsnSerAsn 323
                                                                                                                                                                                                                                                               344 ArgValAspGlyArgArgArgLeuValTyrLysPheGlyLysAsnSerSerGlyTrpLysGlu 363
284 IleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGluGlyValPhe 303
                                                                                                                                                                      324 MetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGluIleLeuGlu 343
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Mismatches:
Indels:
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Matches:
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APPLICATION NUMBER: US/09/020,956
FILING DATE: 09-FEB-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210121.427C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 98104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (206) 622-4900
TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
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519.50
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39.03%
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TYPE: nucleic acid
STRANDEDNESS: single
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CITY: Seattle
STATE: WA
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Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  130 -----SerSerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGly 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          148 MetalapheginglualaLeuaspProglyProPheaspGlnGlySerProPhealaGln 167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   188 AlaProSerProGlySerSerAspValSerThrAlaGlyThrGlyAlaSerArgSerSer 207
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37
73
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               CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNDER: US/09/030,607
FILING DATE: 25-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative:
Mismatches:
Indels:
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Matches:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAC, (206) 682-6031
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
                                                                                                                                        COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                         CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
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519.50
50.97%
39.03%
26.24%
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TYPE: nucleic acid
STRANDEDNESS: single
NUMBER OF SEQUENCES: 224
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Best Local Similarity:
Query Match:
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                                                                       Seattle
                                                                                                            USA
                                                                                                                          98104
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                                                                                                          COUNTRY:
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276 TrpGluPheileArgAspileLeulleHisProGluLeuAsnGluGlyLeuMetLysTrp 295
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         296 GluasnargHisGluGlyValPheLysPheLeuargSerGluAlaValAlaGlnLeuTrp 315
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                                                                                                                                                                                                                                                                        472 CCTGTTGCAGAGTCACCTGATATGAAAAGGAGCAAGACCCCCCCTGCCAAGTGCCACACC 413
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208 HisserSerAspSerGlyGlySerAspValAspLeuAspProThrAspGlyLySLeuPhe 227
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jounifer L.
APPLICANT: Mitcham, Jounifer L.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Farlocker, Susan Louise
APPLICANT: Ralos, Michael
APPLICANT: Ralos, Michael
APPLICANT: Ranger, Gary
APPLICANT: Ranger, Mark
APPLICANT: Renger, Mark
APPLICANT: Bolk, John
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
TITLE OF INVENTION: UNMERR: US/09/439,313
CURRENT PRILICAND DAYE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
IENGTH: 852
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Matches:
Conservative:
                                                                                                             228 ProSerAspGlyPheArgAspCysLysLys-----
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, Patent No. 6329505
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LOCATION: (1)...(852)
OTHER INFORMATION: n = A,T,C or G
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ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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296 GluAsnArgHisGluGlyValPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrp 315
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                                                                                                                                   790 CCTTTCCANGAGTTCGACATCAACGGCGAGCACCTTTGCAGCATGAGTTTGCAGGAGTTC 731
                                                                                                                                                                                                                                                                  111 ArgleuValPheGlyProLeuGlyAspGlnLeuHisAlaGlnLeuArgAspLeuThr--- 129
                                                                                                                                                                                                                                                                                        ----SerSerSerSerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGly 147
                                                                                                                                                                                                                                                                                                                                                                                                                     148 MetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySerProPheAlaGln 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                       649 ---CTGTTCCAGTCC----ACACAC 632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             168 GluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHisProGlySerCysGlyAlaGly 187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          208 HisSerSerAspSerGlyGlySerAspValAspLeuAspProThrAspGlyLysLeuPhe 227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     565 TTATATGACACCAACTATGGTAGCACAGTÁGATTTG-----TTGGACAGCAAAACTTTC 512
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      238 -----GlyAspProLysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLys 255
                                                                                                              71 GlnValLeuAspTrpIleSerTyrGlnValGluLysAsnLysTyrAspAlaSerAlaIle 90
                                                                                                                                                                                                                                                                                                                                                                              670 AACGGCCAGTGCAGTAGTGAC------
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Mismatches:
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                                                                        US-08-978-217-2 (1-371) x US-09-439-313-44 (1-852)
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US-09-352-616A-44/c
; Sequence 44, Application US/09352616A
Best Local Similarity: 39.03%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 649 ---CTGTTCCAGTCC-----ACACAC 632
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                 GENERAL INFORMATION:

APPLICANT: Dillon, Davin C.

APPLICANT: Harlocke Susan Louise

APPLICANT: Harlocke Susan Louise

APPLICANT: Jiang, Yuqui

APPLICANT: Xu, Jiangchun

APPLICANT: Micham, Jennifer Lynn

APPLICANT: Micham, Jennifer Lynn

APPLICANT: Micham, Jennifer Lynn

TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE

TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE

CURRENT APPLICATION NUMBER: US/09/352,616A

CURRENT FILING DATE: 1999-07-13

NUMBER OF SEQ IN NOS: 472

COMMUNER OF SEQ IN NOS: 472
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Matches:
Conservative:
Mismatches:
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OTHER INFORMATION: n = A,T,C or G
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519.50
50.97%
39.03%
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                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapien
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Patent No. 6395278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Percent Similarity:
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US-09-352-616A-44
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ò	256 (256 GluTyrīrpAspCysLeuGluGlyLysLysSerlysHisAlaProArgGlyThrHisLeu 275
đ	412 /	AAAAAGCACAACCGGAGGGACTCACTTA 383
δλ	276	TrpGluPhelleArgAspIleLeulleHisProGluLeuAsnGluGlyLeuMetLysTrp 295
qq	382	AGACAAGAACCCAGGATTA
ò	296	GluasnargHisGluGlyValPheLysPheLeuArgSerGlualaValAlaGlnLeuTrp 315
qq	322	GATCTGAGGGCGTCT
ò	316	316 GlyGlnLysLysLysAsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyr 335
đ	262	GGTAAAAAGAACAACAACAACAGCCATGACCTATGAAAAGCTCAGCGGAGCTATGAGATAT 203
ζŏ	336	TyrTyrLysArgGluIleLeuGluArgValAapGlyArgArgLeuValTyrLysPheGly 355
q	202	
δλ	356	SerGlyTrr
qq	142	
Search Job ti	.complet	Search completed: February 12, 2004, 21:49:55 Job time : 105.522 secs

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February 12, 2004, 21:46:16 ; Search time 64.1925 Seconds (without alignments) 7673.534 Million cell updates/sec
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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1116
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Perfect score:
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/packfiles1.seq:* Issued Patents NA:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Sequence 27, Appl	4,	Н	282	282,	28	282,	282,	44, 4	44,	44,	44	44	'n	'n		92	33	æ	ω̈	<u>۽</u>	'n	۲,	ď	12	Sequence 1, Appli	'n,
	ID	.09	US-09-570-593-4	US-08-746-789A-1	US-09-389-681-282	US-09-620-405B-282	US-09-339-338-282	US-09-433-826B-282	ó	-09-020-956-4		US-09-439-313-44	-09-352-	-60-	-09-009-913-	US-09-009-913-3	US-09-009-913-4	-010-60-	US-09-009-913-338	-09-009-913	-600-60-	-09-009-913-	US-09-213-767-1	US-09-570-593-1	US-09-055-113-2	US-09-570-593-12	-08-36	US-08-368-281-3
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* Query	Match	100.0	100.0	99.4	44.7	44.7	44.7	44.7	44.7	15.6	15.6	15.6	٠ س	'n.	ď.	15.6	15.6	14.9	14.6	9.1	9.1	9.1	7.6	7.0	•	•	9.9	9.9
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Sequence 307, App	Sequence 3, Appli	Sequence 10, Appl	ä	Seguence 1, Appli	'n	840	3, A	Sequence 1, Appli	٦,	Sequence 1, Appli	m	ı,	m	255	17	86	r,
US-09-016-434-307	US-09-920-759-3	US-09-920-759-10	US-09-360-779-1	US-09-435-335-1	US-09-344-579-1	US-09-016-434-840	US-08-343-443B-3	US-08-469-412A-1	US-09-021-715-1	US-08-875-944B-1	US-09-116-049-3	US-09-602-868A-1	US-09-884-363-3	US-09-300-958A-25	US-09-920-759-11	US-09-620-312D-869	US-08-878-177-1
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6.2	6.1	6.1	6.0	6.0	5.9	5.9	5.7	5.4	5.4	5.3	5.3	5.3	5.3	5.2	5.2	5.2	4. 8.
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ALIGNMENTS

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## Sequence 27, Application US/09300958A

## Sequence 27, Application US/09300958A

## Patent No. 6495319

## GENERAL INFORMATION:

## APPLICANT: Mcclelland, Michael

## APPLICANT: Mcclelland, Michael

## APPLICANT: Mcclelland, Michael

## APPLICANT: Tenkle, Thomas

## TITLE OF INVENTION: Reduced Complexity Nucleic Acid Targets and Methods of

## TITLE OF INVENTION: Using Same

## TITLE OF INVENTION: Using Same

## TITLE OF INVENTION: Using Same

## TITLE OF INVENTION: Using Same

## TITLE OF INVENTION: USing Same

## TITLE OF INVENTION: USing Same

## TITLE OF INVENTION NUMBER: 60/083,331

## PRIOR FILING DATE: 1998-04-27

## PRIOR FILING DATE: 1998-08-27

## PRIOR FILING DATE: 1998-08-27

## PRIOR FILING DATE: 1998-02-04

## NUMBER OF SEQ ID NOS: 85

## SEGFWARE: PatentIn Ver. 2.0
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Best Local Similarity 100.0%; Pred. No. 1.6e-273;
Matches 1116; Conservative 0; Mismatches 0; Indels 0;
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RESULT 1
US-09-300-958A-27
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                                                                                                                                                                        LOCATION: (96)...(1211)
OTHER INFORMATION: Human epithelial-restricted with serine box (ESX)
OTHER INFORMATION: protein.
                                                                                                                                                                                                                                                                             100.0%; Score 1116; DB 4; Length 1907; 100.0%; Pred. No. 1.6e-273; tive 0; Mismatches 0; Indels 0;
  SEQ ID NOS: 13
FastSEQ for Windows Version 4.0
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Matches 1116; Conservative
                                                                                     TYPE: DNA ORGANISM: Homo sapiens PEATURE:
NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
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APPLICANT: Xin, Hong
APPLICANT: Xin, Hong
APPLICANT: Harrowe, Greg
APPLICANT: Harrowe, Greg
TITLE OF INVENTION: EXPRESSION OF ETS-DOMAIN PROTEINS IN
FILLE REFERENCE: 2300-1556
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CURRENT FILING DATE: 2000-05-12
PRIOR APPLICATION NUMBER: 60/134,112
PRIOR FILING DATE: 1999-05-14
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Patent No. 6566063
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                       ATCCTGGAACGCGTGGATGGCCGGCGACTCGTCTACAAGTTTGGCAAAAACTCAAGCGGC 1080
                                                                                                                                                                                                                                                                                                                                   1116 ATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACAAGTTTGGCAAAAACTCAAGCGGC 1175
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                    841 GACATCCTCATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAA 900
                                                               936 GACATCCTCATCCACCCGGAGCTCAACGAGGGCCTCATGAAGTGGGAGAATCGGCATGAA 995
                                                                                                                  GGCGTCTTCAAGTTCCTGCGCTCCGAGGCTGTGGCCCAACTATGGGGCCAAAAGAAAAA 960
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/08746789A

Patent No. 5789200
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: A No. 5789200el Human ETS Family Member, ELF3
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 99.4%; Score 1109.6; DB 1; Length 1920; Best Local Similarity 99.6%; Pred. No. 6.7e-272; Matches 1112; Conservative 0; Mismatches 4; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                      1081 TGGAAGGAAGAAGATTCTCCAGAGTCGGAACTGA 1116
                                                                                                                                                                                                                                                                                                                                                                                                                                  1176 TGGAAGGAAGAAGTTCTCCAGAGTCGGAACTGA 1211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: MICROSOFT WORD
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NOBER: US/08/746,789A
No. 5789200ember 15, 1996
N: 514
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TELEPHONE: 610 270 5219
TELEPAX: 610 270 4026
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             34,344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: NO. 57892
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: William T. Han
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 1920
TYPE: Nucleic Acid
STRANDEDNESS: Singl
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HANTI-SENSE: NO
US-08-746-789A-1
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US-08-746-789A-1
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            Sequence 282, Application US/09389681A

Sequence 282, Application US/09389681A

Betent No. 6518237

GENERAL INFORMATION:
APPLICANT: Viqui, Jiang

APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: M, Jiangchun

TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND

TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 21011.470C3

CURRENT APPLICATION NUMBER: US/09/389,681A

CURRENT FILING DATE: 1999-09-02

NUMBER OF SEQ ID NOS: 463

SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                             Similarity 99.8%; Pred. No. 1.8e-117; 009; Conservative 0; Mismatches 1; Indels 0
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Best Local Similarity 99.84
Matches 500; Conservative
                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
COCANISM: Homo sapiens
US-09-389-681-282
US-09-389-681-282
                                                                                                                                                                                                                                                                                            SEQ ID NO 282
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Sequence 282, Application US/09620405B Patent No. 6528054

RESULT 5 US-09-620-405B-282

GENERAL INFORMATION:
APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
APPLICANT: Harlocker, Susan L.
APPLICANT: Hepler, William T.

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CATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACTACAAACGGGAGATCCTGGA 1028
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APPLICANT: Yugiu, Jiang
APPLICANT: Yugiu, Jang
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: LIGHTON DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILLE REPERENCE: 210121.47062
CURRENT APPLICATION NUMBER: US/09/339,338A
CURRENT APPLICATION NUMBER: US/09/339,338A
NUMBER OF SEQ ID NOS: 315
SOFTWARE: FastSEQ for Windows Version 3.0
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAFY AND TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER FILE REFERENCE: 210121.47008
CURRENT APPLICATION WUMBER: US/09/620,405B
CURRENT FILING DATE: 2000-07-20
NUMBER OF SEQ ID NOS: 495
SOFTWARE: FASISEQ for Windows Version 3.0
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Matches 500; Conservative
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US-09-620-405B-282
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US-09-339-338-282
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LENGTH: 502
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                                            609 ITCTCGGAGCTCCCACTCCTCAGACTCCGGTGGAAGTGACGT
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CORGANISM: Homo sapiens
US-09-604-287A-282
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REPERRICE: 210121.470C4
CURRENT APPLICATION NUMBER: US/09/433,826B
CURRENT FILING DATE: 1999-11-03
MUMBER OF SEQ IN NOS: 474-11-03
MUMBER OF SEQ IN NOS: 474-11-03
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Best Local Similarity 99.8%; Pred. No. 1.8e-117;
Matches 500; Conservative 0; Mismatches 1; Indels 0;
                                                            Query Match 44.7%; Score 499.4; DB 4; Length 502; Best Local Similarity 99.8%; Pred. No. 1.8e-117; Matches 500; Conservative 0; Mismatches 1; Indels 0;
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; Patent No. 6579973
; GENERAL INFORMATION:
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CRGANISM: Homo mapiens
US-09-433-826B-282
ORGANISM: Homo sapiens
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    ; US-09-339-338-282
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LENGIH: 502
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                                   729 GAAGCGGAAACGAGGCCGGCCCCGAAAGCTGAGCAAAGAGTACTGGGACTGTCTCGAGGG 788
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APPLICANT: Jiang, Yuqiu
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Mitchan, Jennifer L.
APPLICANT: Mulliam T.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
TILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: COMPOSITIONS OF BREAST CANCER
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
TITLE OF INVENTION: 470C7
CURRENT FILLING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FastsEQ for Windows Version 3.0
SEQ ID NO 282
LENGTH: 502
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GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C,
TITLE OF INVENTION: COMPOUNDS FOR INMUNOTHERAPY OF PROSTATE CANCER AND METHODS
NUMBER OF SEQUENCES: 224
CORRESPONDENCES: SEED and BERRY LLP
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                     998 TGAGGTACTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACA 1057
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                          450 ridaaaaaagdaaagaacccccccccaagridccacaaaaaagcacaaaccccaaaaagaga 391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/030,607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
15.6%; Score 173.8; DB 3;
Best Local Similarity 69.7%; Pred. No. 6.5e-35;
Matches 235; Conservative 0; Mismatches 102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 25-FEB-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: MAKA, David J.
REGISTRATION NUMBER: 31,392
REPERROCE/DOCKET NUMBER: 210121.427C3
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UMBER: US/09/030,607
25-FEB-1998
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-030-607-44/c
; Sequence 44, Application US/09030607
; Patent No. 6262245
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INFORMATION FOR SEQ ID NO: 44:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 98104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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Patent No. 6261562

GENERAL INFORMATION:
APPLICANT: Dilin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                 362 CAAGTTCCTGCGCTCCGAGGCTGTGGCCCAACTATGGGGCCAAAAAGAAAAGAACAGCAA 421
122 TGGCAAGCTCTTCCCCAGCGATGGTTTTCGTGACTGCAAGAAGGGGGGATCCCAAGCACGG 181
                                             729 GAAGCGGAAACGAGGCCGGCCCCGAAAGCTGACAAAGAGTACTGGGGACTGTCTCGAGGG
                                                                                      182 GAAGCGGAAACGAGGCCGGCCCGGAAAGCTGAGCAAAGAGTACTGGGACTGTCTCGAGGG
                                                                                                                                                                                                                           849 CATCCACCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAAGGCGTCTT
                                                                                                                                                                                                                                                          302 CATCCACCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAAGGCGTCTT
                                                                                                                                    789 CAAGAAGAGCAAGCACGCCCAGAGGCACCCACCTGTGGGAGTTCATCCGGGACATCCT
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COUNTRY: USA

COMPUTER: READALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CORPUTER: PREFER PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/020,956
FILING DATE: 09-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
15.6%; Score 173.8; DB 3;
Best Local Similarity 69.7%; Pred. No. 6.5e-35;
Matches 235; Conservative 0; Mismatches 102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: SEED and BERRY LLP 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/ACENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REPERENCE/DOCKET NUMBER: 210121.427C2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1029 ACGGGTGGATGGCCGGCGACT 1049
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INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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STATE:
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998 IGAGGIACTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACA 1057
                                                        270 AGCTATGGGGTAAAAAGAAGAACAACAGCAGCATGACCTATGAAAAAGCTCAGCCGAGCTA 211
                                                                                                                                                                 938 AACTATGGGGCCAAAAGAAAAGAACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCA 997
                                                                                                                                                                                                                  1058 AGTITGGCAAAACTCAAGCGGCTGGAAGGAGGAAGA 1094
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US-09-232-149A-44/c
; Sequence 44, Application US/09232149A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: miso_feature
LOCATION: (1)...(852)
OTHER INFORMATION: n = A,T,C or G
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                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION: APPLICANT: Dillon,
                                                                                                                                                                                                                                                                                                                                     RESULT 12
US-09-352-616A-44/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-352-616A-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 44
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450 rehanandehechaenececeerrecenerecenenenenanneenenenenenenenen 391
                                                                                              CCCACCTGTGGGAGTTCATCCGGGACATCCTCATCCACCGGAGCTCAACGAGGCCTCA 877
                                                                                                                                     390 crcacriardedaarrcarcecedacarccrcrigaaccaadacaagaaccaddarraa 331
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Sequence 44, Application US/09439313
Sequence 44, Application US/09439313
Sequence 41, Application US/09439313
Sequence 42, Application
Sequence 43, Application
Sequence 44, Application
Sequence 47, Subsequence
Septicant: Dilon, Davin C.
Septicant: Harlocker, Subsequence
APPLICANT: Harlocker, Subsequence
APPLICANT: Ralos, Michael
APPLICANT: Ralos, Michael
APPLICANT: Ralos, Michael
APPLICANT: Relos, Michael
APPLICANT: Bolk, John
Solk, John
Septicant: Day, Craig
Septicant: Day, Craig
Septicant: Day, Craig
Septicant: Day, Craig
Septicant: Day, Craig
Septicant: Day, Craig
Septicant: Day, Craig
Septicant: Dilon, Diamosts OF PROSTATE CANCER
FILE REFERENCE: 210121.42709
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT APPLICATION NUMBER: US/09/439,313
SUFFWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; NAME/KEY: misc_feature
; LCCATION: (1)...(852)
; CTHER INFORMATION: n = A,T,C or G
US-09-439-313-44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             818 CCCACCTGTGGGAGTTCATCCGGGACATCCTCATCCACCCGGAGCTCAACGAGGCCTCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
APPLICANT: Dillon, Davin C.
APPLICANT: Barlocker, Susan Louise
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
CURRENT APPLICATION NUMBER: US/99/352,616A
CURRENT FILING DATE: 1999-07-13
NOMBER OF SEQ ID NOS: 472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 15.6%; Score 173.8; DB 4; Length 852; Best Local Similarity 69.7%; Pred. No. 6.5e-35; Matches 235; Conservative 0; Mismatches 102; Indels 0;
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                      GENERAL INFORMATION:
APPLICANT: X1, Jiangchun
APPLICANT: X2, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 21012.42706
CURRENT APPLICATION NUMBER: US/09/232,149A
CURRENT FILING DATE: 1999-01-15
NUMBER OF SEQ ID NOS: 338
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1058 AGTITGGCAAAACTCAAGCGCCTGGAAGGAGGAGA 1094
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COMPUTER READBLE FORM:
COMPUTER READBLE FORM:
COMPUTER: IBM COMPATIBLE
COMPUTER: IEM COMPATIBLE
COMPUTER: FESTERO
COPERATING SYSTEM:
SOFTWARE: FESTERO for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,913
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Patent No. 6087485
GENERAL INFORMATION:
APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
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STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature

LOCATION: (1)...(852)

// CTHER INFORMATION: n = A,T,C or G

US-09-232-149A-44
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ORGANISM: Homo sapien
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Patent No. 6465611
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US-09-009-913-2
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998 TGAGGTACTACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACA 1057
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATURG SYSTEM: DOS
SOFTWARE: FastSEC for Windows Version 2.0
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APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Genes
TIMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
CITY: Palo Alto
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APPLICATION NUMBER: US/09/009,913
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                                                                                                                                                                                 REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: SEC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
FILING DATE: 21-JAN-1998
                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                             TELEFAX: 650-327-3231
                    CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: CDNA
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US-09-009-913-3
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STATE:
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Db 1092 AATTTGGGAAGAATGCCCGAGGATGGAGGAAATGA 1128

1058 AGTITGGCAAAACTCAAGCGGCTGGAAGGAGGAAGA 1094

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Search completed: February 13, 2004, 01:54:40 Job time : 65.1925 secs

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APPLICANT: Sebastian HORRSCH
APPLICANT: Sebastian HORRSCH
APPLICANT: Shubhangi KANATKAR
APPLICANT: Shubhangi KANATKAR
APPLICANT: Shubhangi KANATKAR
APPLICANT: Reter OLANDT
APPLICANT: Michael MORRISEY
APPLICANT: Michael MORRISEY
APPLICANT: APPLICANT: APET VIEW
APPLICANT: APPLICANT: Rosematie SCHWANDT
APPLICANT: Rosematie SCHWANDT
APPLICANT: Karen LU
APPLICANT: Karen LU
APPLICANT: Karen GLATT
APPLICANT: Kanen GLATT
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MRI-030
                       Sequence 101, App
Sequence 363, App
Sequence 192, App
Sequence 1105, Ap
Sequence 1105, Ap
Sequence 217, App
Sequence 64, App
Sequence 853, App
Sequence 853, App
Sequence 863, App
Sequence 284, App
Sequence 282, App
Sequence 282, App
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Sequence 44, Appl
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Sequence 203, App
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Sequence 1953, Ap
Sequence 12, Appl
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Sequence 44,
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5 US-10-097-340-74
0 US-09-964-824A-101
0 US-09-964-824A-101
0 US-09-880-107-3420
0 US-09-880-107-3420
0 US-09-922-217-1105
2 US-10-025-380-1105
2 US-10-025-380-1105
2 US-10-025-380-3853
0 US-09-922-217-9853
0 US-09-922-217-9853
0 US-09-933-263-944
0 US-10-025-380-944
0 US-09-933-263-944
0 US-09-938-2290
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US-09-030-605-44

US-09-030-605-44

US-09-115-453-44

US-09-115-453-44

US-09-115-453-44

US-09-115-453-44

US-09-115-453-44

US-09-895-814-44

US-09-895-814-44

US-10-12-896-44

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US-10-012-94-025-44

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: John MONAHAN
APPLICANT: MAJULIA GANNAVARAPU
APPLICANT: Sebastian HOGRSCH
APPLICANT: Shubhangi KAWATKAR
APPLICANT: Steve G. KOVATS
APPLICANT: Rachel B. MEYERS
APPLICANT: Michael MORRISEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 4 9 9 1 1 4 4 2 6 9 1 4 4 2 6 6 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 8 5 2 8 5 2 8 5 2 8 5 2 8 5 2 8 5 2 8 5 2 8 5 2 8 5 2 8 5 2 
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-MODEL=frame+ p2n.model -DEV=x1h
-MODEL=frame+ p2n.model -DEV=x1h
-Q0/cgn2 1/USFTO_spool/US08978217/runat_10022004_133827_20514/app_query.fasta_1.1500
-Q2/cgn2 1/USFTO_spool/US08978217/runat_1002204_133827_20514/app_query.fasta_1.1500
-DB=Published_Applications_NA -QFMT=fastap -SUPFTX=mpb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR SCORE=pct -THR MAX=100
-THR MIN=0 -ALIGN=15 -MODE=LGCAL -OUTFYT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXLEN=2000000000 -USES=US08978217 occn 1 1 690 @runat 1002204 133827_20514
-NCPU=6 -ICPU=3 -NO MMAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPBLOCK=100
-LONGLOG -DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                  February 12, 2004, 21:34:51 ; Search time 583.817 Seconds (without alignments) 2334.540 Million cell updates/sec
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| cgn2_6/ptodata/1/pubpna/USO7_PUBCOMB.seq:*
| cgn2_6/ptodata/1/pubpna/USO7_NEW PUB.seq:*
| cgn2_6/ptodata/1/pubpna/USO6_NEW PUB.seq:*
| cgn2_6/ptodata/1/pubpna/USO6_PUBCOMB.seq:*
| cgn2_6/ptodata/1/pubpna/USO6_PUBCOMB.seq:*
| cgn2_6/ptodata/1/pubpna/USO8_PUBCOMB.seq:*
                            GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                      - nucleic search, using frame_plus_p2n model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2449703 segs, 1841816367 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                      US-08-978-217-16_COPY_2_371
                                                                                                                                                                                                                                                                                                                                                                                            Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Maximum DB seq length: 200000000
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Score Match Length DB
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Perfect score:
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                                                                                                                                                                       Run on:
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No
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200 ThralaThrProGlnSerSerHisAlaSerAspSerGlyGlySerAspValAspLeuAsp
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US-09-964-824A-101
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Mismatches:
Indels:
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Matches:
CURRENT APPLICATION NUMBER: US/10/097,340
PRIOR FILING DATE: 2002-03-14
PRIOR PLING DATE: 2002-03-14
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-6
PRIOR PLING DATE: 2001-03-6
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-14
PRIOR PLING DATE: 2001-03-16
PRIOR PLING DATE: 2001-09-26
PRIOR PLING DATE: 2001-09-10
PRIOR PLING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-10
PRIOR PRIOR PRIOR APPLICATION NUMBER: 60/325,102
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-19
PRIOR PRILING DATE: 2001-09-19
PRIOR PRILING DATE: 2001-09-19
SOFTWARR: FREUSEQ for Windows Version 4.0
SECIENGTH: 1907
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92.72%
86.79%
86.21%
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ORGANISM: Homo sapiens
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GENERAL INFORMATION:
APPLICANT: HOCKIGAN, Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signation: TITLE OF INVENTION: cates
TITLE OF INVENTION: cates
TITLE OF INVENTION: cates
TITLE OF INVENTION: dets
TITLE OF INVENTION: dets
TITLE OF INVENTION: dets
TITLE OF INVENTION: dets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/60/236,033
PRIOR PAPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
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PRIOR APPLICATION NUMBER: US/60/236,028
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                       755
                                                                            239
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220 LeuThrGluSerLygValPheProArgAspAspAspPheThrAspTyrLygLysGlyGluPro
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US-09-964-824A-101
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Best Local Similarity:
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Db 1200 TGGAAGGAGGAAGGTTCTCCAGAGTCGGAAC 1232	RESULT 3 US-964-824A-563 ; Sequence 563, Application US/09964824A ; Patent No. US20020102531A1	; GENERAL INFORMATION: ; APPLICANT: Horrigan, Stephen ; TITLE OF INVENTION: Cancer Gene Determination and T ; TITLE OF INVENTION: Sets	; FILE REFERENCE: 689290-73 ; CURRENT APPLICATION NUMBER: US/09/964,824A ; CURRENT FILING DATE: 2001-09-27 ; FRIOR APPLICATION NUMBER: US/60/236,033	; PRIOR PILING DATE: 2000-09-28 ; PRIOR APPLICATION NUMBER: US/60/236,032 ; PRIOR FILING DATE: 2000-09-28 ; PRIOR APPLICATION NUMBER: US/60/236,028	; PRIOR FILING DATE: 2000-09-28 ; NUMBER OF SEQ ID NOS: 583 ; SEQ ID NO 563	; LENGTH: 1915 ; TYPE: DNA ; ORGANISM: Homo sapiens US-09-964-824A-563	1.19e-189 Length: 1707.00 Matches:	Percent Similarity: 92.72\$ Conservative: Best Local Similarity: 86.79\$ Mismatches: Query Match: 86.21\$ Indels: DB: 10	US-08-978-217-16_COPY_2_371 (1-370) x US-09-964-824A-5 QY 1 AlaAlaThrCygGluIleSerAsnValPheSerAsnTyrP	Db 123 GCTGCAACCTGTGAGATTAGCAACATTTTAGCAACTACTACT	183 GAGACTCCACCCTGGCTCTGTTCCCCCTGC	Oy 40 LeuThrLeuAsnAsnGlnGlnMetThrLeuGluGlyProG	60 GluargproGlnPheTrpSerLyaTh :::	80 LysAsnLysTyrAspAlaSerSerIleAspPheSer	Oy 100 LeuCysSercysAlaLeuGluGluLeuArgLeuValPheG :::	Qy 120 HisAlaGlnLeuArgAspLeuThrSerAsnSerSerAspG	140 LeuLeuGluLysAspGlyMetSerPhe	Db 543 CTGCTGGAGAAGGATGGCCATGGCCTTCCAGGAGGCCCTA- Oy 160 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyA	Db 600 CAGGCCACCCTTTGCCCAGGAGCTGCTGGACGGTG
	21 GluasproThrLeualaBroAlaBroProThrThrPheGlyThrGluaspLeuVal 39	40 LeuThrLeuAsnAsnGinGlinMetThrLeuGluGlyProGluLysAlaSerTrpThrSer 59	60 GluargProGlnFheTrpSerLysThrGlnValLeuGluTrpIleSerTyrGlnValGlu 79	80 LysasnlysTyraspalaSerSerIleaspPheSerargcysasnMetaspGlyalathr 99 	100 LeuCysSerCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeu 119	120 HisaladinLeuArgAspLeuThrSerAsnSerSerAspGluLeuSerTrpIleIleGlu 139	140 LeuleuGluLysAspGlyMetSerPheGlnGluSerLeuGlyAspLeuGlyProPheAsp 159	160 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyArgGlnAlaSerProTyrTyr 179	180 CysserThrTyrGlyProGlyAlaProSerProGlySerSerAspValSerThrAlaArg 199 :::	200 ThralaThrProGlnSerSerHisAlaSerAspSerGlyGlySerAspValAspLeuAsp 219	LeuThrGluSerLysValPheProArgAspAspPheThrAspTyrLysLysGlyGluPro		AAGCACGGGAAGCGGAAACGAGGCCGGCCGGAAAGCTGAAGCAAGAAGAGTACTGGGAACTGT LEUGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheileArg			1020 GGCGTCTTCAAGTTCCTGCGCTCCGAGGGTGTGGCCCAACTATGGGGCCAAAGAAAAG 1079 320 AsnSerAsnMetThrTyrGlulysLeuserArgAlaMetArgTyrTyrTyrTysArgGlu 339	AACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAG 11	340 IleteuGluArgyalAspGlyArgArgLeuValTyrLysPheGlyLysAsanSerSerGly 359 1140 ATCCTGGAACGGCTGGATGGCCTGACGTCTACAAGTTTGGCAAAAACTCAACGGC 1199	360 TrpLysGluGluGluValGlyGluSerArgAsn 370
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APPLICANT: Horne, Darci T.
APPLICANT: Wockley, Joseph G.
APPLICANT: Generf, Uwe
APPLICANT: Gener Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer;
FILE REPERBUCE: 44921-5028-W0
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT PLING DATE: 2001-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950

ACTIVATE: PARCEL PARCELLIN VENT OF A PRIOR FILING DATE: 200-10-02
NUMBER OF SEQ ID NOS: 3950
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                                                        300 GlyValPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLys 319
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                             | 160 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyArgGlnAlaSerProTyr 179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: COMPOUNDS FOR IMPUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: COMPOUNDS FOR IMPUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE FILE REPRENCE: 210121.471C13
CURRENT APPLICATION NUMBER: 0.09/9922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTMARE: FRESEQ FOR Windows Version 4.0
SEQ ID NO 1105
LENGTH: 1917
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Regher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
Clapper, Jonathan D.
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Sequence 192, Application US/09967768A

Patent No. US20021050877A1

SEQUENCE 192, Application US/09967768A

Patent No. US20021050877A1

SENERAL INPORMATION:

APPLICANT: Augustus, Meena

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu

TITLE OF INVENTION: Sets

FILE REPRENCE: 689290-72

CURRENT PILING DATE: 2001-09-28

PRIOR APPLICATION NUMBER: US/60/236,109

PRIOR PILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 325

SOUTHARE: PatentIn version 3.0

SEQ ID NO 192

LENGTH: 1915
LeuLeuGlulysAspGlyMetSerPheGlnGluSerLeuGlyAspLeuGlyProPheAsp 159
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                                      Length:
Matches:
Conservative:
Mismatches:
Indels:
                                      1.19e-189
1707.00
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US-09-922-217-1105
                           Alignment Scores:
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1022 GGCGTCTTCAAGTTCCTGCGCTCCGAGGCTGTGGCCCCAACTATGGGGCCAAAAGAAAAG 1081
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                                                                        320 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGlu
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USF
FILE REFERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFFWARE: FASUSED for Windows Version 4.0
SEQ ID NO 1105
LENGTH: 1917
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APPLICANT: Lodes, Michael J.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Reagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Smith, Carole L.
King, Gordon E.
Wang, Aljun
Clapper, Jonathan D.
Skeiky, Yasir A. W.
Fanger, Gary R.
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US-10-025-380-1105
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1082 AACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGG 1141
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                                          LysAsnLysTyrAspAlaSerSerIleAspPheSerArgCysAsnMetAspGlyAlaThr 99
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PAL33P1
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/18569
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Publication No. US20040005579A1
GENERAL INFORMATION:
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Mismatches:
Indels:
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Matches:
PRIOR FILING DATE: 2001-06-07
PRIOR PELLON NUMBER: US 60/209,467
PRIOR FILING DATE: 2000-06-07
NUMBER OF SEQ ID NOS: 4360
SEPTAMER: Patentin Ver. 3.1
SEQ ID NO 756
LENGTH: 1956
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                                                                                                                                  TYPE: DNA ORGANISM: Homo sapiens
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HisAlaGlnLeuArgAspLeuThrSerAsnSerSerAspGluLeuSerTrpIleIleGlu 139
                                                             504 CATGCCCAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAG
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Publication No. US20030235915A1
GENERAL INFORMATION:
APPLICANT: SPECHT, THOMAS
APPLICANT: SCHWITT, ARMIN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: POSENTHAL, ANDRE
APPLICANT: POSENTHAL, ANDRE
TITLE OF INVENTION: TUMAN NUCLEIC ACID SEQUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMANS
FILE REFERENCE: SCH-1763
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US-10-131-410-64
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                                                                                                                       1001 GACATICTICATCCACCCGGAGCTYAACGAGGGCCTYATGAAGTGGGGAGAATCGGCATGAA 1060
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                                                                                                                                                                       300 GlyValPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLys 319
                                                                                       280 AspileLeuileHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 299
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  260 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg
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TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REPERENCE: PA106
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR PILING DATE: 2000-03-08
PRIOR PILING DATE: 1999-03-12
NUMBER: OF ROSEN OF PRIOR PILING DATE: 1999-03-12
NUMBER: OF SEQ ID NOS: 1694
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 207
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Matches:
Conservative:
Mismatches:
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92.72%
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86.21%
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; ORGANISM: Homo sapiens
US-09-925-301-207
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Best Local Similarity:
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US-09-925-301-207
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Mismatches:
                                                                                                                                                                                                                                                                                     Length:
Matches:
CURRENT APPLICATION NUMBER: US/10/131,410
CURRENT FILING DATE: 2002-04-25
FRIOR APPLICATION NUMBER: 09/646,673
PRIOR FILING DATE: 2000-09-20
PRIOR PILING DATE: 1990-09-19
NUMBER OF SEQ ID NOS: 202
SOFTWARE: Patentin Ver: 2.1
SOFTWARE: 2269
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1458.50
93.71%
88.36%
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; ORGANISM: Homo sapiens
US-10-131-410-64
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Best Local Similarity:
Query Match:
DB:
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Pred. No.:
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313 LeuTrpGlyGlnLysLysLysAsnSerAsnMetThrTyrGluLysLeuSerArgAlaMet 332
                                                                                                                                                                              790 crardegeccaaaaaaaaaaaaacaecaacarcarcaacaagaagcraagccecarg 849
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; Patent No. US20020076414A1
, GENERAL INFORMATION:
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US-09-922-217-853
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US-09-922-217-853/c
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US-08-978-217-16_COPY_2_371 (1-370) x US-10-025-380-853 (1-626)
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Secrist, Heather
Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
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Skeiky, Yasir A. W.
Fanger, Gary R.
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
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US-10-025-380-853/c
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                                     324 GCCCTA---GACCCAGGGCCCTTTGACCAGGCCAGCCCCTTTGCCCCAGGAGCTGCTGGAC
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384 TCTGATGAGCTCAGTTGGATCATTGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAG 325
                                                                                                                                                                                                                                                                                           147 TCCGGTGGAAGTGAACGTGGAACCTGGATCCCACTGATGGCAAGCTCTTCCCCAGCGATGGT 88
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Stolik, John
APPLICANT: Stolik, John
APPLICANT: Stolik, John
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471012
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT APPLICATION NUMBER: US/09/833,263
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                        171 AspGlyArgGlnAlaSerProTyrTyrCysSerThrTyrGlyProGlyAlaProSerPro
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Patent No. US20020110547A1
GENERAL INFORMATION:
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131 SerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGlyMetSerPheGlnGlu 150
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51 GlyProGluLysAlaSerTrpThrSerGluArgProGlnPheTrpSerLysThrGlnVal
                                                                                  71 LeuGluTrpIleSerTyrGlnValGluLysAsnLysTyrAspAlaSerSerIleAspPhe
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CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 944
LENGTH: 563
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Patent No. US20020076414A1
GENERAL INFORMATION:
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
Clapper, Jonathan D.
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US-09-922-217-944
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Sequence 944, Application US/09833263
| Patent No. US2020110547A1
| GENERAL INFORMATION:
| APPLICANT: Wang, Aljun
| APPLICANT: Glapper, John Au
| APPLICANT: Stolk, John Au
| APPLICANT: Stolk, John Au
| APPLICANT: COMPOUNDS FOR INMUNOTHERAPY AND
| TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
| TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
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Search completed: February 13, 2004, 01:46:24 Job time : 599.817 secs

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Sequence 74, Appl Sequence 101, App Sequence 563, App Sequence 192, App Sequence 1105, App Sequence 756, App Sequence 207, App Sequence 207, App Sequence 64, App Sequence 1661, App Sequence 1661, App Sequence 1953, App Sequence 1953, App
                                                                                                                                         February 13, 2004, 01:27:22 ; Search time 2732.91 Seconds (without alignments) 10448.744 Million cell updates/sec
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-036-2379 Sequence 2319, Ap 5-98-2216 Sequence 2216, Ap 6-036-226 Sequence 2216, Ap 5-88-229 Sequence 2220, Ap 5-874-282 Sequence 282, App 6-05-282 Sequence 323, App 6-05-284 Sequence 944, App 6-05-284 Sequence 243, App 6-05-284 Sequence 243, App 6-05-284 Sequence 253, App 6-05-284 Sequence 253, App 6-05-284 Sequence 253, App 6-05-284 Sequence 253, App 6-05-284 Sequence 12739, A 6-117-253 Sequence 12739, A 6-117-253 Sequence 12739, A 6-117-253 Sequence 253, App 1-05-253, App 1-05	es and Proteins For The Identification, and Therapy of Ovarian Cancer
15 US-10-060-036-22 16 US-09-998-598-522 17 US-09-604-287A-28 18 US-09-604-287A-28 19 US-09-604-287A-28 11 US-09-551-621-28 13 US-10-076-652-28 14 US-10-076-652-28 15 US-10-076-652-28 16 US-09-998-598-32 17 US-09-998-32 18 US-10-085-117-28 18 US-10-085-117-25 19 US-09-864-761-1476 10 US-09-864-761-1476 11 US-09-972-546-16 12 US-10-085-117-25 13 US-10-085-117-25 14 US-09-864-761-1476 15 US-10-085-117-25 16 US-09-864-761-1476 17 US-09-864-761-1476 18 US-10-085-117-25 19 US-09-864-761-1476 11 US-09-972-546-16 12 US-10-085-117-523-71 12 US-10-085-117-224 13 US-10-085-117-224 14 US-10-085-117-28	ALIGNMENTS 10097340 10097340 10197340 1021 1032 10332 103324,967 106/325,102 106/325,102 106/325,102 106/325,102 106/325,102
22.2 22.1 22.1 22.1 22.1 23.1 24.2 25.1 26.2 27.1 27.1 27.2	RESULT 1 US-10-097-340-74 Sequence 74, Application US/10097340 PUDICATION NO. US20030087250A1 GENERAL INFORMATION: APPLICANT: JOHN MONHAN APPLICANT: School GENCRACH APPLICANT: Steve GENCRATARR APPLICANT: Steve GENCRATARR APPLICANT: Steve GENCRATARR APPLICANT: Steve GENCRATARR APPLICANT: Steve GENCRATARR APPLICANT: Steve GENCRATARR APPLICANT: Rachel E. MEYERS APPLICANT: Rachel E. MEYERS APPLICANT: Rachel E. MEYERS APPLICANT: Rachel E. MEYERS APPLICANT: Rachel E. MEYERS APPLICANT: Rachel E. MEYERS APPLICANT: Rachel D. MILLS APPLICANT: Rachel D. MILLS APPLICANT: Resement GENCRATARR APPLICANT: Resement GENCRATARR APPLICANT: Resement, Dr. APPLICANT: Resement, Prefitte OF INVENTION: NUCLEIC ACID MOI TITLE OF INVENTION: ASSESSMENT, PRIOR FILING DATE: 2001-03-14 PRIOR PLING DATE: 2001-03-14 PRIOR PLING DATE: 2001-03-14 PRIOR PLING DATE: 2001-03-14 PRIOR PLING DATE: 2001/09/26 PRIOR PLING DATE: 2001/09/26 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16 PRIOR PLING DATE: 2001-03-16
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; Sequence 563, Application US/09964824A
; Patent No. US20020102531A1
; GENERAL INFORMATION:
      CURRENT FILING DATE: 2001-09-27
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Patent No. US20020102531A1
APPLICANT: HORINATION: Lephen
TITLE OF INVENTION: Sets
FILE OF INVENTION: Sets
FILE REPERRACE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
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                                                                                                                                                                                               Query Match 2.8%; Score 216; DB 15; Length 1907;
Best Local Similarity 66.7%; Pred. No. 3.3e-54;
Matches 456; Conservative 0; Mismatches 195; Indels 33; Gaps
PRIOR APPLICATION NUMBER: 60/323,580 PRIOR FILING DATE: 2001-09-19 NUMBER OF SEQ ID NOS: 363 SOFTWARE: PastSEQ for Windows Version 4.0
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                                                                                                                      ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-097-340-74
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US-09-964-824A-101
                                                            SOFTWARN. SEQ ID NO 74
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PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 101
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; ORGANISM: Homo sapiens
US-09-964-824A-101
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APPLICANT: Horrigan, Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signaturities of INVENTION: Sets
FILE OF INVENTION: Sets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT APPLICATION NUMBER: US/60/236,033
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
SOFTWARE: Patentin version 3.0
SEQ ID NO 563
LENGTH: 1915.
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; ORGANISM: Homo sapiens
US-09-964-824A-563
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                                                                                                                                APPLICANT: Horne, Darci T.

APPLICANT: Horne, Darci T.

APPLICANT: Wockley, Joseph G.

APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.

TILE OF INVENTION: Gene Expression Profiles in Liver Cancer;
FILE REFRENCE: 44921-5028-WO
CURRENT APPLICATION WINGHER: US/09/880,107

CURRENT APPLICATION NUMBER: US/09/880,107

CURRENT PILING DATE: 2000-06-14

PRIOR FILING DATE: 2000-06-14

PRIOR FILING DATE: 2000-06-14

PRIOR FILING DATE: 2000-10-02

NUMBER OF SEQ ID NOS: 3950

SOFTWARE PATENTIN VET. 2.1
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US-09-880-107-3420
                           ; Sequence 3420, Application US/09880107; Patent No. US20020142981A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
US-09-880-107-3420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 3420
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1656 CACCTCCCTGCTCAGTGCTTGGGCTCCACGGGCAGGGGTCAGAGGACTCCCTAATTTATG 1715
                                                                                                                                                                                                  1716 İĞÜTATATAATATÇEÇAĞATĞTAÇATAĞAĞATÇTATTTTTTTTTAAAÇATTCCCTCCC 1775
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65.6%; Pred. No. 4.4e-54;
tive 0; Mismatches 204; Indels 31; Gaps
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APPLICANT: Lodes, Michael J.
APPLICANT: Lodes, Michael J.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Wang, Tongrong
APPLICANT: Wing, Yuqiu
APPLICANT: Sining, Yuqiu
APPLICANT: Ming, Aljun
APPLICANT: Ming, Aljun
APPLICANT: Clapper, Jonathan D.
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1105
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CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
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Patent No. US20020076414A1
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Best Local Similarity 65.6%
Matches 449; Conservative
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COGANISM: Homo sapiens
US-09-922-217-1105
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| Sequence 192, Application US/09967768A
| Patent No. US20020150877A1
| GENERAL INFORMATION:
| APPLICANT Augustus, Meena
| ITLE OF INVENTION: Sets
| FILE REFERENCE: 689280-7.
| ITLE OF INVENTION: Sets
| FILE REFERENCE: 689280-7.
| FILE REPERENCE: 689280-7.
| FILE REPERENCE: 689280-7.
| FILE OF INVENTION NUMBER: US/60/236,109
| PRIOR PILIOR DATE: 2000-09-28
| PRIOR FILING DATE: 2000-09-28
| PRIOR FILING DATE: 2000-09-28
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                 6999 AGGIATITACIACAAACGGGAGAICCIGGAACGGGIGGAIGGCCGACGGCTCGICIACAAG
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65.6%; Pred. No. 4.4e-54;
tive 0; Mismatches 204; Indels 31; Gaps
                                                                                                 7631 CTGCTCTTCTCCACTGAGTGCTGG 7654
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Matches 449; Conservative
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ORGANISM: Homo sapiens
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US-09-967-768A-192
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US-09-967-768A-192
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7398 ATGAGACCCACCTGCAGATGCCTGGCCTCAGCCAAGGAGGCTGGGGAGACTGTGGCAGGA 7457
                                                                  7458 GACTGCAGGGACGGACAGGGTTGTGTCCTCGG----TACTTCCTGGACTGCCTTC 7513
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65.6%; Pred. No. 4.4e-54;
tive 0; Mismatches 204; Indels 31; Gaps
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FREESE for Windows Version 4.0
                                                                                                                                                                                                                                                                             7631 CTGCTCTTCTCCACTGAGTGCTGG 7654
                                                                                                                                                                                                                                                                                                              1778 CACTCCTCTCCCACAGAGTGCTGG 1801
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Clapper, Jonathan D.
Skeiky, Yasir A. W.
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Smith, Carole L.
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Best Local Similarity 65.64
Matches 449; Conservative
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; ORGANISM: Homo sapiens
US-10-025-380-1105
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LENGTH: 1917
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1301 GGACAGGCCAGATGG-CCCCTCCACTGGGGAATGCTCCCAGCTGTGGTGGAGAG 1359
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                                                                                                                         7239 GAAGAACCTGTTGGGCGTGCCCTCTGC---AGTCTCCTCAAGTGCAGCCTTTGGCCTC--
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65.6%; Pred. No. 4.4e-54;
tive 0; Mismatches 204; Indels 31; Gaps
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Sequence 756, Application US/10264049

Bublication No. US20040005579A1

GENERAL INFORMATION:
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PAJ39P1
CURRENT FILING DATE: 2002-10-04

PRIOR FILING DATE: 2001-06-07

PRIOR FILING DATE: 2001-06-07

PRIOR FILING DATE: 2001-06-07

PRIOR FILING DATE: 2000-06-07

PRIOR FILING DATE: 2000-06-07

SOFTWARE: PACENTIN NUMBER: US 60/209,467

NUMBER OF SEQ ID NOS: 4360

SOFTWARE: PACENTIN VEF. 3.1
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Best Local Similarity
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6999 AGGIAITACTACAAACGGGAGATCCTGGAACGGGTGGATGGCCGACGGCTCGTCTACAAG 7058
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CORGANISM: Homo sapiens
US-10-131-410-64
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LENGTH: 2269
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                                                                                      7239 GAAGAACCIGITGGGCGIGCCCICIGC --- AGICICCICAAGIGCAGCCITTGGCCTC 7293
                                                                                                                               1399 AAGCTGATGTTTTGGTGTATTGTCAGCCATCGTCCTGGGACTCGGAGACTATGGCCTCGC 1458
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Pred. No. 4.5e-54;
0; Mismatches 204; Indels 31; Gaps
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TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REPERENCE: PA106
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PATENTIN VET: 2.0
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Patent No. US20020052308A1
GENERAL INFORMATION:
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Best Local Similarity 65.6%;
Matches 449; Conservative (
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ORGANISM: Homo sapiens
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Best Local Similarity 65.6%; Pred. No. 4.9e-54;
Matches 449; Conservative 0; Mismatches 204; Indels 31; Gaps
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APPLICANT: SPECHT, THOMAS
APPLICANT: SCHMITT, ARMIN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: ROSSWTHAL, ANDRE
APPLICANT: ROSSWTHAL, ANDRE
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMAN SECUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMANS
FILE REPERBNCE: SCH-1763
CURRENT APPLICATION NUMBER: US/10/131,410
CURRENT FILING DATE: 2002-04-25
PRIOR FILING DATE: 2000-09-20
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6421 ATGACCTATGAGAAGCTGAGCCGAGCCATGAGGTGAG 6457
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OTHER INFORMATION: a, t, c, g, unknown or other
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OTHER INFORMATION: a, t, c, g, unknown or other
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LOCATION: (1145). (1145)
OTHER INFEMATION: a, t, c, g, unknown or other
US-10-22-798-1601
                                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY: modified_base
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LOCATION: (201)..(1235)
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ORGANISM: Homo sapiens
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Publication No. US203035833A1
Sequence 1601, Application US/10292798
Publication No. US2030235833A1
SEDERAL INFORMATION:
APPLICANT: SURA, MAKIKO
APPLICANT: ASAI, KIYOSHI
APPLICANT: ALWARA, TUTAKA
APPLICANT: ALWARATANA, HIROYUKI
TITLE OF INVENTION: GUANOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS
FILE REPERENCE: 084335/166
CURRENT APPLICATION NUMBER: US/10/292,798
CURRENT PILING DATE: 2002-11-13
PRIOR PILING DATE: 2001-12-18
PRIOR FILING DATE: 2001-12-18
PRIOR PILING DATE: 2001-12-18
PRIOR FILING DATE: 2001-106-18
NUMBER OF SEQ ID NOS: 2070
SOSTWARR PACENTION VOLES.
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850 AGGTACTACTACAAACGGGAATCCTGGAACGGGTGGATGGCCGGGACTCGTCTACAAG 909
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ORGANISM: Homo sapiens
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LENGTH: 1435
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                                                 0; Gaps
Query Match 2.4%; Score 185.8; DB 12; Length 1435; Best Local Similarity 79.4%; Pred. No. 4.7e-45; Matches 220; Conservative 0; Mismatches 57; Indels 0;
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RESULT 15
US-10-025-380-853/c
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LENGTH: 626
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                                                                                     Query Match
2.4%; Score 185.8; DB 13; Length 1435;
Best Local Similarity 79.4%; Pred. No. 4.7e-45;
Matches 220; Conservative. 0; Mismatches 57; Indels 0;
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
PILE REPERBECE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
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; LOCATION: (1145)
; OTHER INFORMATION: a, t, c, g, unknown or other US-10-017-161-1953
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Masqher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 853, Application US/09922217
Patent No. US20020076414A1
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Smith, Carole Lynn
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Wang, Aijun
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US-09-922-217-853
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Best Local 9
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APPLICANT:
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4267 CCGCTGCAACATGGACGGAGCCACCTCTGCAGCTGTGCGCTGGAGGAGCTGCGGCTAGT 4326
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502 ACGATGTGACATGGATGGCGCCACCCTCTGCAATTGTGCCCCTTGAGGAGCTGCGTCTGGT 443
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Stolik, John A.
APPLICANT: Stolik, John A.
APPLICANT: Stolik, John A.
APPLICANT: Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILLS REPERENCE: 210121.471012
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT APPLICATION NUMBER: US/09/833,263
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FRRESCO FOR Windows Version 3.0
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2.3%; Score 176.2; DB 10; Length 626;
Best Local Similarity 85.6%; Pred. No. 2.3e-42;
Matches 196; Conservative 0; Mismatches 33; Indels 0;
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; Publication No. US20020182191A1
                                                                                                                                                                                                                                                                                           Sequence 853, Application US/09833263
Patent No. US20020110547A1
GENERAL INFORMATION:
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Stolk, John A.
Wang, Tongtong
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Vedvick Thomas S.
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
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Smith, Carole L.
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Wang, Aijun
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CRGANISM: Homo sapien
US-09-833-263-853
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; FILE REFERENCE: 210121.471C14
; CURRENT APPLICATION WUMBER: US/10/025,380
; CURRENT FILING DATE: 2001-12-19
; NUMBER OF SEQ ID NOS: 1129
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 853
LENGTH: 626
; TYPE: DNA
; ORGANISM: HOMO Sapiens
US-10-025-380-853
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Sequence 927, App Sequence 2290, App Sequence 2282, App Sequence 282, App Sequence 163, App Sequence 163, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 192, App Sequence 1105, App
                                                                                                                                     February 13, 2004, 01:27:22; Search time 7.40341 Seconds (without alignments) 10448.744 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpna/USO7_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/USO7_PUBCOMB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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0 US-09-998-598-32

US-09-998-598-2290

US-09-604-287A-282

1 US-09-639-338-282

1 US-09-551-621-282

3 US-10-007-805-282

5 US-10-007-805-282

5 US-10-007-805-282

6 US-10-964-824A-101

0 US-09-964-824A-563

0 US-09-964-824A-563

0 US-09-967-768A-192

0 US-09-967-768A-192

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0 US-09-967-768A-192
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                                                                                                                                                                                                                                                                                                                                                                                                     2449703 seqs, 1841816367 residues
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Maximum Match 1008
Listing first 45 summaries
                                                                                        OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                  1 GTACCTCATGGCCCGGGTCAG 21
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Maximum DB seq length: 200000000
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21
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Match Length DB
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Perfect score:
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                                                                                                                                                                                                                                                                                  Sequence:
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                                                                                                                                     Run on:
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Sequence 1105, Ap Sequence 756, App Sequence 207, App Sequence 3261, App Sequence 1501, Ap Sequence 1501, Ap Sequence 1955, App Sequence 149050, Sequence 562, App Sequence 599, App Sequence 201, App Sequence 10965, Sequence 25, App Sequence 10, App Sequence 10, App Sequence 1139, Ap Sequence 201, App Sequence 201, App Sequence 201, App Sequence 201, App Sequence 201, App Sequence 201, App Sequence 201, App Sequence 201, App Sequence 201, App Sequence 201, App Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 65, Appli Sequence 65, Appli Sequence 65, Appli
US-10-025-380-1105 US-10-0264-049-756 US-10-922-301-301 US-10-131-410-64 US-10-106-036-3261 US-10-106-036-3261 US-10-1056-761-11 US-10-1056-761-11 US-10-1056-761-11 US-10-107-632-149050 US-10-027-632-149050 US-10-027-632-149050 US-10-027-632-149050 US-10-027-632-149050 US-10-10-027-632-109965 US-10-10-027-632-109965 US-10-10-027-632-109965 US-10-10-027-632-109965 US-10-10-027-632-109965 US-10-10-027-632-109965 US-10-10-027-632-109965 US-10-10-10-10-10-10-10-10-10-10-10-10-10-
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1917 1956 1996 1996 1996 1996 1996 1435 1435 1435 1935 1935 1930 1930 1930 1930 1930 1930 1930 1930
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11111222222222222222222222222222222222
00000 000 0 000

ALIGNMENTS

RESULT 1 US-10-305-720-927/c

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Sequence 927, Application US/10305720

Publication No. US20040010136A1

GENERAL INFORMATION:
TITLE OF INVENTION: Composition for the Detection of Signaling Pathway Gene Expressiving TITLE OF INVENTION: COMPOSITION FOR THE 2002-11-26

FILE REFERENCE: PA-0002-1 CON
CURRENT APPLICATION NUMBER: US/10/305,720

CURRENT APPLICATION NUMBER: 09/016,434

PRIOR FILING DATE: 1998-01-30

FRIOR FILING DATE: 1998-01-30

SOFTWARE: PERL Program
SEQ ID NO 927

LENGTH: 237

TYPE: DNA
ORGANISM: Homo sapiens
FRATURE:
FRATURE:
FRATURE:
OTHER INFORMATION: Incyte ID No. US20040010136A1 773734

US-10-305-720-927

Query Match
Best Local Similarity 100.0%; Pred. No. 0.77;
Matches 21; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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TYPE: DNA
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                                                                                                                        ; ORGANISM: Homo sapiens
US-09-604-287A-282
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US-09-339-338-282/c
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US-09-551-621-282/c
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                                                                                                                         TYPE: DNA
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APPLICANT: Dillon, Javin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Matcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
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APPLICANT: Susan L.
APPLICANT: Susan L
                      APPLICANT: Chenall, Ruth A. APPLICANT: Chenall, Ruth A. APPLICANT: Meagher, Madelein Joy TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER FILE REFERENCE: 210121.561
CURRENT APPLICATION NUMBER: US/09/998,598
CURRENT PILING DATE: 2001-11-16
NUMBER: OF SEQ ID NOS: 2606
SOFTWARE: Corixa Invention Disclosure Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AND METHODS FOR THE THERAPY AND
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100.0%; Score 21; DB 10; Length 451;
Best Local Similarity 100.0%; Pred. No. 0.73;
Matches 21; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2290, Application US/09998598
Patent No. USZ00201150922A1
GENERAL INFORMATION:
APPLICANT: Stolk, John A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Chenault, Ruth A.
APPLICANT: Meagher, Madelein Joy
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER FILE REFERENCE: 210121.561
CURRENT APPLICATION NUMBER: US/09/998,598
CURRENT APPLICATION NUMBER: 2001-11-16
NUMBER OF SEQ ID NOS: 2606
SOFTWARE: COCIXA INVENTION Disclosure Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 282, Application US/09604287A; Patent No. US20020064872A1; GENERAL INFORMATION;
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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Xu, Jiangchun
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CORGANISM: Homo sapiens
US-09-998-598-32
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; ORGANISM: Homo sapiens
US-09-998-598.2290
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US-09-604-287A-282/c
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US-09-998-598-2290
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LENGTH: 451
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Sequence 282, Application US/0933933A
; Sequence 282, Application US/0933933A
; Patent No. US20020102602A1
; GENBEAL INFORMATION:
    APPLICANT: Widiu, Jaing
; APPLICANT: Dillon, Davin C.
    APPLICANT: Mitcham, Jennifer L.
    APPLICANT: Mitcham
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
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APPLICANT: Harlocker, Susan L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210121.470CS
CURRENT APPLICATION NUMBER: US/09/551,621
CURRENT FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 4779
SOFTWARE: FastSEQ for Windows Version 3.0
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100.0%; Score 21; DB 9; Length 502;
Best Local Similarity 100.0%; Pred. No. 0.72;
Matches 21; Conservative 0; Mismatches 0; Indels
SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 282 LENGTH: 502
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Best Local Similarity 100.0
Matches 21; Conservative
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RESULT 9
US-10-076-622-282/c
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APPLICANT: Dillon, Davin C.
APPLICANT: Dillon, Davin C.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Herlocker, Susan L.
APPLICANT: Henderson, Robert A.
APPLICANT: Fanger, Gary R.
APPLICANT: Penger, Gary R.
APPLICANT: Penger, Gary R.
APPLICANT: Penger, Gary R.
APPLICANT: OWNOALS, Thomas S.
APPLICANT: Durham, Margarita D.
APPLICANT: MCNoill, Patricia D.
APPLICANT: MONOILL, Patricia D.
APPLICANT: MONOILL, AND DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.470C10
CURRENT APPLICATION NUMBER: US/10/007,805
CURRENT FILING DATE: 2001-12-07
NUMBER OF SEQ ID NOS: 593
SOFTWARE FEASTER FEASTER FOR Windows Version 4.0
                                                                                                                                                                  Sequence 282, Application US/10124805
| Publication No. US20030166022A1
| GABREAL INFORMATION:
| APPLICANT: Houghton, Raymond L. APPLICANT: Bleath, Paul R. |
| APPLICANT: Bleath, Paul R. |
| APPLICANT: Persing, David H. |
| TILLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TILLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER FILE REFERENCE: 210121.470C12
| CURRENT FILING DATE: 2002-04-15
| NUMBER OF SEQ ID NOS: 627
| SOFTWARE FEASTE FASTER FOR Windows Version 4.0
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100.0%; Score 21; DB 13; Length 502;
Best Local Similarity 100.0%; Pred. No. 0.72;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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Best Local Similarity 100.0%; Pred. No. 0.72;
Matches 21; Conservative 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 282, Application US/10007805
Publication No. US20020150581A1
GENERAL INFORMATION:
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                                                458 GTACCICATGGCCCCGCTCAG 438
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1 GTACCTCATGGCCCGGCTCAG 21
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ORGANISM: Homo sapiens
US-10-007-805-282
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US-10-124-805-282
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                                                                                                                                               US-10-124-805-282/c
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SEQ ID NO 282
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LENGTH: 502
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TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer FILE REFERENCE: MRI-030
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Sequence 282, Application US/10076622

Sequence 282, Application US/10076622

Publication No. US20030023036A1

GENERAL INFORMATION:

APPLICANT: Sleath, Paul R.

APPLICANT: Sleath, Paul R.

TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER

PILE REPREBRICE 2 210121.47011

CURRENT APPLICATION NUMBER: US/10/076,622

CURRENT APPLICATION NUMBER: US/10/076,622

CURRENT FILING DATE: 2002-02-13

NUMBER OF SEQ ID NOS: 627

SOFTWARE PRESENCE: Anidows Version 4.0

SEQ ID NO 282

LENGTH: 502
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CURENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR PLICATION NUMBER: 60/325,149
PRIOR APPLICATION NUMBER: 60/326,026
PRIOR APPLICATION NUMBER: 60/324,967
PRIOR APPLICATION NUMBER: 60/324,967
PRIOR PILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/314,732
PRIOR PILING DATE: 2001-09-10
PRIOR PLING DATE: 2001-09-10
PRIOR PLING DATE: 2001-09-10
PRIOR PLING DATE: 2001-09-10
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PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SOFTWARE: FASESEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 74, Application US/10097340; Publication No. US20030087250A1; GENERAL INFORMATION: APPLICANT: John MONAHAN; APPLICANT: Sebatian HOERSCH; APPLICANT: Shabhangi KAMATKAR; APPLICANT: Steve G. KOVATS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          458 GTACCTCATGGCCCGGCTCAG 438
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Rosemarie SCHMANDT
Kumei ZHAO
Karen GLATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peter VEIBY
Gordon B. MILLS
Robert C. BAST, Jr.
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Michael MORRISEY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
; ORGANISM: Homo sapiens
US-10-076-622-282
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Sequence 192, Application US/09967768A

| Sequence 192, Application US/09967768A
| Patent No. US2002015087741
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| Application of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of the properties of t
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Patent No. US20020142981A1

Patent No. US20020142981A1

Patent No. US20020142981A1

Patent No. US20020142981A1

APPLICANT: Horentain Darci T.

APPLICANT: Generi, Uwe

APPLICANT: Gene Logic, Inc.

TILE OF INVENITON: Gene Expression Profiles in Liver Cancer: TILE OF INVENITON: Gene Expression Profiles in Liver Cancer: TURRENT APPLICATION NUMBER: US/09/880,107

CURRENT APPLICATION NUMBER: US 60/211,379

PRIOR PLING DATE: 2000-06-14

PRIOR PLING DATE: 2000-06-14

PRIOR PLING DATE: 2000-06-14

PRIOR PLING DATE: 2000-10-02

NUMBER OF SEQ ID NOS: 3950

SOFTWARE PATENTING OF US CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO CONDO 
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US-09-880-107-3420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 21; DB 10; Length 1915; Pred. No. 0.65;
                                   100.0%; Score 21; DB 10; Length 1915; 100.0%; Pred. No. 0.65; 0; Mismatches 0; Indels 0.
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Best Local Similarity 100.0%; Pred. No. 0.65;
Matches 21; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                 1124 Graccreárescecescreas 1104
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Matches 21; Conservative
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US-09-967-768A-192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
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Best Local Similarity
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US-09-967-768A-192/c
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US-09-880-107-3420/c
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LENGTH: 1915
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Sequence 563, Application US/09964824A

Patent No. US20020102531A1

GENERAL INFORMATION:

APPLICANT: Horrigan, Stephen

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

FILE REPERENCE: 689290-73

CURRENT APPLICATION NUMBER: US/60/236,033

PRIOR PILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

SOFTWARE: PatentIn version 3.0

SEQ ID NO 563

LENGTH: 1915
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; Patent No. US20020102531A1
; GENERAL INFORMATION:
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
; TITLE OF INVENTION: Sets
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CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT PILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR PILING DATE: 2000-09-28
PRIOR FILING DATE: 5000-09-28
NUMBER OF SEQ ID NOS: 583
SOFTWARE: Patentin version 3.0
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US-09-964-824A-563
                                                                                           ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-097-340-74
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US-09-964-824A-101
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US-09-964-824A-101/c
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US-09-964-824A-563/c
SEQ ID NO 74
LENGTH: 1907
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LENGTH: 1915
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Sequence 1105/Application US/09922217
Sequence 1105, Application US/09922217
Sequence 1105, Application US/09922217
SEXEAL INFORMATION:
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Sitclk, John A.
APPLICANT: Sitclk, John A.
APPLICANT: Sinth, Carole Lynn
APPLICANT: Mang, Yuqiu Gordon E.
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09/922,217
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1105
LENGTH: 1917
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100.0%; Score 21; DB 9; Length 1917;
Best Local Similarity 100.0%; Pred. No. 0.65;
Matches 21; Conservative 0; Mismatches 0; Indels C
  0; Mismatches
                                             Matches 21; Conservative
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ORGANISM: Homo sapiens
US-09-922-217-1105
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US-09-922-217-1105/c
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Search completed: February 13, 2004, 11:48:34 Job time : 16.4034 secs

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Sequence 282, App. Sequence 1601, Ap Sequence 1601, Ap Sequence 74, Appl Sequence 101, App Sequence 563, App Sequence 192, App
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Sequence 2290, Ap
Sequence 282, App
Sequence 282, App
Sequence 282, App
Sequence 282, App
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10448.744 Million cell updates/sec
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                                                                                                                                                         February 13, 2004, 01:27:22 ; Search time 7.40341 Seconds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Published Applications NA:*

| cgn2 6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO6 NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO6 NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO6_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO6_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO8_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO8_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO8_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO8_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO9_NEW_PUB.seq:*
| cgn2 6/ptodata/1/pubpna/USO0_NEW_PUB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-064-228A-220

US-09-339-338-2282

US-09-339-338-282

US-09-339-338-282

US-00-339-338-282

US-10-124-885-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-282

US-10-07-805-81

US-09-964-824A-101

US-09-984-8563

US-09-9880-107-3420
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Maximum Match 100%
Listing first 45 summaries
                                                                                                           OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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Match Length DB
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APPLICANT: Kales, Michael D.
APPLICANT: Lodge, Michael J.
APPLICANT: Lodge, Michael J.
APPLICANT: Deraing, David H.
APPLICANT: Peraing, David H.
APPLICANT: Hepler, William T.
APPLICANT: Hepler, William T.
APPLICANT: Jiang, Yudiu T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER FILE REFERENCE: 210121.566
CURRENT APPLICATION NUMBER: US/10/060,036
CURRENT FILING DATE: 2002-01-30
NUMBER OF SEQ ID NOS: 4560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 21; DB 15; Length 275; 100.0%; Pred. No. 2.5;
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LENGTH: 275
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; Patent No. US20020150922A1
; GENERAL INFORMATION:
US-10-060-036-3261
; Sequence 3261, Application US/10060036
; Publication No. US20030073144A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 109 CCGGGACATCCTCATCCACCC 129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
CORGANISM: Homo sapiens
US-10-060-036-3261
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Best Local Similarity
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US-09-998-598-2290/c
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Sequence 282, Application US/09339338A

Patent No. US20020102602A1

GENERAL INFORMATION:
APPLICANT: Vigiu, Jiang
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: MITCHAM, DIANOSING OF BREAST CANCER AND
TITLE OF INVENTION: DIANOSING OF BREAST CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.470C2
GURRENT APPLICATION UNBERS. US/09/339,338A

CURRENT PILING DATE: 1999-06-23
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: AU, Jiangchun
APPLICANT: Harlocker, Susan L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
FILE REPERENCE: 210121.470C7
CURRENT PILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 488
SOFTWARE: FastSEQ for Windows Version 3.0
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                  APPLICANT: XL, Jiangchun
APPLICANT: Chenault, Ruth A.
APPLICANT: Meagher, Madelein Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
CURRENT APPLICATION NUMBER: US/09/998,598
CURRENT PILING DATE: 2001-11-16
NUMBER OF SEQ ID NOS: 2606
SOFTWARE: Corixa Invention Disclosure Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
100.0%; Score 21; DB 10; Length 499;
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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APPLICANT: Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ORGANISM: Homo sapiens
US-09-604-287A-282
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US-09-998-598-2290
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US-09-339-338-282
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LENGTH: 499
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SNGTH: 502
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APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
APPLICANT: Micham, Jennifer L.
TITLE OF INVENTION: OWNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE FILE REPERENCE: 210121.470C5
CURRENT APPLICATION NUMBER: US/09/551,621
CURRENT APPLICATION NUMBER: US/09/551,621
NUMBER OF SEQ ID NOS: 4779
SOFTWARE: FastSEQ for Windows Version 3.0
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100.0%; Score 21; DB 13; Length 502;
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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100.0%; Score 21; DB 11; Length 502;
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels C
                                                                                                                                                                                                           Query Match 100.0%; Score 21; DB 10; Length 502; Best Local Similarity 100.0%; Pred. No. 2.4; Matches 21; Conservative 0; Mismatches 0; Indels 0
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APPLICANT: Sleath, Paul R.
APPLICANT: Sleath, Paul R.
APPLICANT: Persing, David H.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.470C12
CURRENT APPLICATION NUMBER: US/10/124,805
CURRENT FILING DATE: 2002-04-15
SOFTWARE: FASES OF FOR WINDOWS VERSION 4.0
                    SOTTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 282 LENGTH: 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 282, Application US/10124805; Publication No. US20030166022A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 5
US-09-551-621-282
Sequence 282, Application US/09551621
Publication No. US20030104366A1
GENERAL INFORMATION:
APPLICANT: Yuqui, Jiang
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NUMBER OF SEQ ID NOS: 315
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ORGANISM: Homo sapiens
US-10-124-805-282
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; ORGANISM: Homo sapiens
US-09-551-621-282
                                                                                                           TYPE: DNA
CRGANISM: Homo sapiens
US-09-339-338-282
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US-10-124-805-282
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Page 3

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Sequence 1601, Application US/10292798

Sequence 1601, Application No. US2003235833A1

Septicant: SUMA, MAKIXO

SAPELCANT: SUMA, MAKIXO

SAPELCANT: ASUA, MAKIXO

SAPELCANT: ASUA, MINOSHI

APPLICANT: ASUA, MINOSHI

APPLICANT: ASUA, MINOSHI

TITLE OF INVENTON: GUANOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS

FILE REFERENCE: 084335/166

CURRENT APPLICATION NUMBER: US/10/292,798

CURRENT FILING DATE: 2002-11-13

PRIOR PILING DATE: 2001-12-18

RIOR PILING DATE: 2001-06-18

PRIOR PILING DATE: 2001-06-18

PRIOR PILING DATE: 2001-06-18

NUMBER OF SEQ ID NOS: 2070

SOFTWARE: Patentin Ver: 2.1

SEQ ID NO 1601

LENGTH: 1435

LENGTH: 1435
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Publication No. US20030143668A1

GENERAL INFORMATION:

APPLICANT: SUMA, MAKINO

APPLICANT: AKIYAMA, YUTAKA

APPLICANT: AKIYAMA, YUTAKA

APPLICANT: ARIXAMA, YUTAKA

FILLE OF INVENTON NOVEL G PROTEIN-COUPLED RECEPTORS

FILE REFERENCE: 084335/0152

CURRENT PILING DATE: 2002-12-18

FRIOR FILING DATE: 2001-06-18

PRIOR FILING DATE: 2001-06-18

PRIOR FILING DATE: 2001-06-18

NUMBER OF SEQ ID NOS: 2430

SOFTWARE PETENTIN VET. 2.1

SEQ ID NO 1953

LENGTH 1435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (1040)..(1139)
OTHER INFORMATION: a, t, c, g, unknown or other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; LOCATION: (1145)..(1145); OTHER INFORMATION: a, t, c, g, unknown or other US-10-292-798-1601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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Best Local Similarity 100.0
Matches 21; Conservative
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LOCATION: (201)..(1235)
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FEATURE:
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; Publication No. US20030023036A1
; GENERAL INFORMATION:
APPLICANT: Houghton, Raymond L.
APPLICANT: Blacth, Paul R.
APPLICANT: Pershing, David H.
ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
ITLE OF INVENTION: AND DIAGMOSIS OF BREAST CANCER
FILE REPERROGE: 210121.47011
CURRENT FAPLICATION NUMBER: US/10/076,622
CURRENT FILING DAVE: 2002-02-13
NUMBER OF SEG ID NOS: 627
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 502
                                                                                                                                                                                                                                                                                APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Marcler, Susan L.
APPLICANT: Hardereon, Robert A.
APPLICANT: Hendereon, Robert A.
APPLICANT: Hendereon, Robert A.
APPLICANT: Panger, Gary R.
APPLICANT: Powdick, Thomas S.
APPLICANT: Wordeil, Patricia D.
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
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APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
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APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: Durham, Margarita
APPLICANT: WONGILL A.0000
APPLICANT: MURGANG: 210121.470C10
CURRENT FILING DATE: 2001-12-07
MUMBER OF SEQ ID NOS: 593
ANDER OF MARKE: PastSEQ for Windows Version 4.0
APPLICANT: SO2
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100.0%; Score 21; DB 14; Length 502;
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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                                                     290 ccecedacarcercarceacec 310
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; ORGANISM: Homo sapiens
US-10-076-622-282
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CRGANISM: Homo sapiens
US-10-007-805-282
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                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
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                                                                                                                         NAME/KEY: modified base
LOCATION: (1040)..(1139)
OTHER INFORMATION: a, t, c, g, unknown or other
                                                                                                                                                                                                                                                       LOCATION: (1145)
COTHER INFORMATION: a, t, c, g, unknown or other US-10-017-161-1953
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PRIOR PELLING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-03-14
PRIOR PILING DATE: 2001-03-14
PRIOR PILING DATE: 2001-09-26
PRIOR PELLING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-10
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PRIOR PELLING DATE: 2001-09-19
PRIOR PELLING DATE: 2001-09-19
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PRIOR PELLING DATE: 2001-09-19
PRIOR PELLING DATE: 2001-09-19
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Publication No. US20030087250A1
GENERAL INFORMATION:
APPLICANT: John MONHAN
APPLICANT: Manjula GANRAVARAFU
APPLICANT: Subsatian HOERSCH
APPLICANT: Steve G. KOVATS
APPLICANT: Steve G. KOVATS
APPLICANT: Michael MORRISEY
APPLICANT: Michael MORRISEY
APPLICANT: Michael MORRISEY
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Robert C. BAST, Jr.
Karen LU
Rosemarie SCHMANDT
                                                                                                                                                                                                                                    NAME/KEY: modified_base
                                                                          LOCATION: (201)..(1235)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peter VEIBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Karen GLATT
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ORGANISM: Homo sapiens
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LOCATION: (1)..(1435)
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Best Local Similarity
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                        FEATURE:
NAME/KEY: CDS
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FATCHER NO. USCUZZULZIALAL

APPLICANT: HOTRIGAN. Stephen

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat

TITLE OF INVENTION: Sets

FILE REFERENCE: 689290-73

CURRENT APPLICATION NUMBER: US/09/964,824A

CURRENT APPLICATION NUMBER: US/09.236,033

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 583

SOFTWARE: PATCHTIN VERSION 3.0

SEQ ID NO 101

LENGTH: 1915
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TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat
TITLE OF INVENTION: Sets
FILE REPERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/60/236,033
PRIOR PILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER: OF SEQ ID NOS: 583
SOFTWARE: PATCHTON NUMBER: US/60/236,028
NUMBER OF SEQ ID NOS: 583
LENGTH: 1915
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0; Mismatches
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                                                                                                                                                                                                            , Sequence 101, Application US/09964824A ; Patent No. US20020102531A1
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                                                                                             932 CCGGGACATCCTCATCCACCC 952
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21; Conservative
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US-09-964-824A-563
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US-09-964-824A-101
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Best Local Similarity
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US-09-964-824A-563
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Sequence 192, Application US/09967768A

Sequence 192, Application US/09967768A

Patent No. US20020150877A1

GENERAL INFORMATION:

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu TITLE OF INVENTION: Set 8

FILE REFERENCE: 689290-72

CURRENT APPLICATION NUMBER: US/09/36,109

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 325

NUMBER OF SEQ ID NOS: 325

SOFTWARE: PETENTION VINCENTION 3.0
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                   Sequence 3420, Application US/09880107
Fatent No. US20020142981A1
GENERAL INFORMATION:
APPLICANT: Horne, Daxci T.
APPLICANT: Wockley, Joseph G.
APPLICANT: Scherf, Usc.
TITLE OF INVENTION: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer:
FILE REFRENCE: 4491-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PATENTIN Ver. 2.1
LENGTH: 1915
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Best Local Similarity 100.0%; Pred. No. 2.3;
Matches 21; Conservative 0; Mismatches 0; Indels 0
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ORGANISM: Homo sapiens
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CORGANISM: Homo sapiens
US-09-967-768A-192
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US-09-967-768A-192
US-09-880-107-3420
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LENGTH: 1915
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Search completed: February 13, 2004, 11:48:25

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Sequence 927, Application US/10305720
Sequence 927, Application No. US20040010136A1
GENERAL INFORMATION:
THILE OF INVENTION: Composition for the Detection of Signaling Pathway Gene Expressic TITLE OF INVENTION: Composition for the Detection of Signaling Pathway Gene Expressic TITLE OF INVENTION: US/10/305,720
CURRENT APPLICATION NUMBER: US/10/305,720
CURRENT PILING DATE: 2002-11-26
PRIOR PRIOR PRING DATE: 1998-01-30
NUMBER OF SEQ ID NOS: 1490
SOGTWARE: PERL PROGRAM
SEQ ID NO 927
LIENGTH: 237
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Sequence 678, App
Sequence 33421, A
Sequence 3, Appli
Sequence 125191,
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Sequence 64, Appl
Sequence 40, Appl
Sequence 1730, Ap
Sequence 3052, Ap
                                                                                                                                         Sequence 74, Appl
Sequence 101, Appl
Sequence 563, App
Sequence 3420, App
Sequence 192, App
                                                           Sequence 2290, Ap
Sequence 3233, Ap
Sequence 2379, Ap
                                                                                                                                                                                                                                                                                     Sequence 1105, Ap
Sequence 1105, Ap
Sequence 756, App
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Sequence 167020,
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Sequence 90917, A
Sequence 4493, Ap
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Sequence 760, App
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; OTHER INFORMATION: Incyte ID No. US20040010136A1 773734
US-10-305-720-927
US-09-190-246-1
US-09-983-965-4018
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           US-10-305-720-927
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46.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Command line parameters:
-MODEL-frame+ p2n.model -DEV=xlh
-Q=CQG12_1/VGPFO_EXPOOL/VGS08978217/runat_10022004_133827_20514/app_query.fasta_1.1500
-Q=CQG12_1/VGPFO_EXPOOL/VGS08978217/runat_10022004_133827_20514/app_query.fasta_1.1500
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-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=A=5 -DOCALATGN=200 -THR SCORE=p5ot -THR MAX-CH=0.1
-THR MIN-0 -ALIGN=15 -MODE=LOCAL -OUTFWT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXIEN=200000000 -USER=US08978217_@CGN 1 1 690 @runat 10022004 133827_20514
-NGPG=6 -LOOPUS - NORM=PT-ARGRQCHERY -NEG SCORE=0 -NATT -DSPBLCCK=100
-LONGLOG -DEV TIMEOUT=120 -WARN TIMEOUT=3 0 -THRANS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                        February 12, 2004, 21:34:51; Search time 25.2461 Seconds (without alignments) 2334.540 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpna/USO7_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/DCT_NEW_PUB.seq:*
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18: /cgn2_6/ptodata/1/pubpna/USO0_NEW_PUB.seq:*
                                         GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                     - nucleic search, using frame_plus_p2n model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2449703 segs, 1841816367 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                               BLOSUM62
Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Maximum DB seq length: 2000000000
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Match Length DB
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                                                                                                                                                                                                                                                                                                                                                    Perfect score:
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                                                                                                                                                                                                                                                                                                                            Title:
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Sequence 2230, Application US/09998598

Sequence 2230, Application US/09998598

Patent No. US20020150922A1

GENERAL INFORMATION:

APPLICANT: Stolk, John A.

APPLICANT: Chemault, Ruth A.

APPLICANT: Chemault, Ruth A.

APPLICANT: Chemault, Ruth A.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: COMPOSITION WORDER: US/09/998,598

CURRENT APPLICATION NUMBER: US/09/998,598

CURRENT FILING DATE: 2001-11-16

NUMBER OF SEQ ID NOS: 2606

SOFTWARE: Corixa Invention Disclosure Database

SEQ ID NO 2230

LENGTH: 499
                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Xu, Jiangchun

APPLICANT: Chanault, Madelein Joy

APPLICANT: Meagner, Madelein Joy

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER

FILE REFERENCE: 210121.561

CURRENT APPLICATION VUMBER: US/09/998,598

CURRENT PILING DATE: 2001-11-16

NUMBER OF SEQ ID NOS: 2606

SOUTHARRE: Corixa Invention Disclosure Database

SEQ ID NO 32

LENGTH: 451
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                                                                                                                                                                                                                                   116 AAAAACTCAAGCGGCTGGAAGGAGGAAGGGTTCTCCAGAGTCGGAAC 163
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Conservative:
Mismatches:
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Mismatches:
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Length:
Matches:
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Gaps:
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                                                                                                                                                                                                                                                                                                                              Sequence 32, Application US/09998598
Patent No. US20020150922A1
GENERAL INFORMATION:
3.23e-06
84.00
100.00%
100.00%
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CORGANISM: Homo sapiens
US-09-998-598-2290
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ORGANISM: Homo sapiens
US-09-998-598-32
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Best Local Similarity:
Query Match:
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Best Local Similarity:
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Pred. No.;
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    Pred. No.:
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APPLICANT: Syle, Ruth A.
APPLICANT: Yu, Jiangchun
APPLICANT: Indirias, Carol Yoseph
APPLICANT: Indirias, Carol Yoseph
APPLICANT: Indirias, Carol Yoseph
APPLICANT: Garter, Heather
APPLICANT: Garter, Darrick
APPLICANT: Garter, Darrick
APPLICANT: Smith, Carole L.
APPLICANT: Sholk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.563
CURRENT APPLICATION NUMBER: US/10/066,543
CURRENT FILING DATE: 2002-01-31
NUMBER OF SEQ ID NOS: 3417
SOFTWARE: FastSEQ for Mindows Version 4.0
SEQ ID NO 3233
LENGTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2379, Application US/10060036

publication No. US20030073144A1

special INFORMATION:
APPLICANT: Benson, Darin R.
APPLICANT: Persing, Michael D.
APPLICANT: Persing, David H.
APPLICANT: Hepler, William T.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER
TITLE TITLE OF INVENTION: TOWN TITLE OF THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER
FILE REFERENCE: 210121.566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63 AAAAACTCAAGCGCCTGGAAGGAGGAAGAGGTTCTCCAGAGTCGGAAC 110
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Conservative:
Mismatches:
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                                                                                                                                            US-08-978-217-12 (1-16) x US-09-998-598-2290 (1-499)
                                                                                                                                                                                                                                                                                                                                 ; Sequence 3233, Application US/10066543; Publication No. US20030087818A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Jiang, Yuqiu
APPLICANT: Pyle, Ruth A.
APPLICANT: Xu, Jiangchun
APPLICANT: Indirias, Carol Yoseph
APPLICANT: Lodes, Michael J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
LOCATION: 438
OTHER INFORMATION: n = A,T,C or G
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84.00
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100.00%
100.00%
Percent Similarity: 100.00% Best Local Similarity: 100.00% Query Match: 100.00% DB:
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Best Local Similarity:
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Sequence 101, Application US/09964824A

Sequence 101, Application US/09964824A

Sequence 101, Application US/09964824A

Sequence 101, Application US/09964824A

Sequence 101, Application US/090102531A1

GENERAL INFORMATION:
TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

TITLE OF INVENTION: Sets

CURRENT FILING DATE: 2000-09-28

SPRIOR PEPLICATION NUMBER: US/60/236,033

PRIOR PELING DATE: 2000-09-28

PRIOR APPLICATION NUMBER: US/60/236,032

PRIOR PELING DATE: 2000-09-28

PRIOR PELING DATE: 2000-09-28

PRIOR PELING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 583

SOFTWARE: Patentin Version 3.0

SEQ ID NO 101

LENGTH: 1915
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Patent No. US20020102531A1
GENERAL INFORMATION:
APPLICANT: HORITION:
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signaturities of INVENTION: Sets
TITLE OF INVENTION: AS Set Sets
FILE REFERENCE: 69290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT PILING DATE: 2001-09-27
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR PILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
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                                                                                         Length:
Matches:
Conservative:
Mismatches:
Indels:
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Matches:
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84.00
100.00%
100.00%
                                                                                            3.02e-05
84.00
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100.00%
; ORGANISM: Homo sapiens
US-10-097-340-74
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APPLICANT: Xumei ZHAO
APPLICANT: Xumei ZHAO
APPLICANT: Xumei ZHAO
APPLICANT: Xumei ZHAO
APPLICANT: Xumei ZHAO
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MR-030
CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
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PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SOFFWARE: FRAESEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              63 AAAAACTCAAGCGGCTGGAAGGAGGAAGAGGTTCTCCAGAGTCGGAAC 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 LysAsnSerSerGlyTrpLysGluGluGluValLeuGlnSerArgAsn 16
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Matches:
Conservative:
Mismatches:
Indels:
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CURRENT APPLICATION NUMBER: US/10/060,036 CURRENT FILING DATE: 2002-01-30 NUMBER OF SEQ ID NOS: 4560 SOPWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2379 LENGTH: 620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 74, Application US/10097340 Publication No. US20030087250A1 GENERAL INFORMATION: APPLICANT: John MONAHAN
                                                                                                                                                                                                       NAME/KEY: misc_feature
) ICCATION: 169, 456, 487, 488, 602
; OTHER INFORMATION: n = A,T,C or G
US-10-060-036-2379
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Sebastian HOERSCH
Shubhangi KAWATKAR
Steve G. KOVATS
Rachel E. MEYERS
Michael MORRISEY
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Robert C. BAST, Jr.
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                                                                                                                                            TYPE: DNA ORGANISM: Homo sapiens
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Best Local Similarity:
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LENGTH: 1907
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DB:
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1187 AAAAACTCAAGGGGCTGGAAGGAGGAAGAGGTTCTCCCAGAGTCGGAAC 1234
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121, 471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
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Matches:
Conservative:
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Conservative:
Mismatches:
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  CURRENT FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: US/60/236,109
PRIOR FILING DATE: 2000-09-28
PRIOR PELICATION NUMBER: US/60/236,034
PRIOR FILING DATE: 2000-09-28
PRIOR PILING DATE: 2000-09-28
PRIOR PILING DATE: 2000-09-28
NUMBER: OF SEQ ID NOS: 325
SOFTWARE: Patentin version 3.0
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SOFTWARE: FastSEQ for Windows Version 4.0
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Patent No. US20020076414A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Xu, Jiangchun APPLICANT: Lodes, Michael J. APPLICANT: Secrist, Heather APPLICANT: Benson, Darin R. APPLICANT: Meagher, Madeleine Joy APPLICANT: Stolk, John A. APPLICANT: Mang, Tongtong
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Homo sapiens
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Best Local Similarity:
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US-09-922-217-1105
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                                                                                                                                                                                                    SEQ ID NO 192
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DB:
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TITLE REFERENCE: 689290-72
CURRENT APPLICATION NUMBER: US/09/967,768A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3420, Application US/09880107

Patent No. US20020142981A1

GENERAL INPORMATION:

APPLICANT: HORNE, Darci T.

APPLICANT: Gene Logic, Inc.

TITLE OF INVENTION:

FILE REPRENCE: 44921-5228-MO

CURRENT FILING DATE: 2001-06-14

PRIOR APPLICATION NUMBER: US 60/211,379

PRIOR APPLICATION NUMBER: US 60/211,379

PRIOR APPLICATION NUMBER: US 60/211,379

PRIOR APPLICATION NUMBER: US 60/217,054

PRIOR PILING DATE: 2000-10-02

NUMBER OF SEQ ID NOS: 3550

SOFTWARE: PATENTIN UND NOS: 2350

SOFTWARE: PATENTIN UND NOS: 2350

SOFTWARE: PATENTIN UND NOS: 2350

SOFTWARE: PATENTIN UND NOS: 2350

SOFTWARE: PATENTIN UND NOS: 2350

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                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Genbank Accession No. US20020142981A1 U73843 US-09-880-107-3420
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Matches:
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Mismatches:
Indels:
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Matches:
Conservative:
Mismatches:
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US-09-967-768A-192
Sequence 192, Application US/09967768A
Patent No. US20020150877A1
GENERAL INFORMATION:
APPLICANT: AUGUSTUS, Meena
NUMBER OF SEQ ID NOS: 583
SOFTWARE: Patentin version 3.0
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                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
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PEATURE:
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Best Local Similarity:
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                                           SEQ ID NO 563
LENGTH: 1915
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TYPE: DNA ORGANISM: Homo sapiens
Percent Similarity:
Best Local Similarity:
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LENGTH: 2269
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                                         Query Match:
DB:
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US-10-131-
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                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Vedvick Thomas S.
APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.41C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1105
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Publication No. US20040005579A1
Publication No. US20040005579A1
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA137P1
CURRENT APPLICATION NUMBER: US/10/264,049
PRIOR APPLICATION NUMBER: PCT/US01/18569
PRIOR FILING DATE: 2001-06-07
PRIOR APPLICATION NUMBER: US 60/209,467
PRIOR APPLICATION NUMBER: US 60/209,467
PRIOR APPLICATION NUMBER: US 60/209,467
NUMBER OF SEO ID NOS: 4360
SOFTWARE: Patentin Ver. 3.1
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Mismatches:
Indels:
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Matches:
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                                  Sequence 1105, Application US/10025380
PUblication No. US20020182191A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Georges, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Mang, Tongtong
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Skeiky, Yasir A. W.
Fanger, Gary R.
Vedvick Thomas S.
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
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ORGANISM: Homo sapiens
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CORGANISM: Homo sapiens
US-10-264-049-756
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Best Local Similarity:
Query Match:
                   -10-025-380-1105
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Pred. No.:
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LENGTH: 1956
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APPLICANT: HINZWANN, BERND
APPLICANT: SCHAITT, ARMIN
APPLICANT: SCHAITT, ARMIN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: DAHL, EDGAR
APPLICANT: DAHL, EDGAR
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUB OF BREAST
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUB OF BREAST
TITLE OF INVENTION: HUMBER: US/10/131,410
CURRENT APPLICATION NUMBER: 09/646,673
FRIOR PILING DATE: 2002-04-25
FRIOR PAPLICATION NUMBER: PCT/DE99/00908
FRIOR PILING DATE: 1999-03-19
FRIOR PILING DATE: 1999-03-19
FRIOR PILING DATE: 1999-03-19
FRIOR PILING DATE: 1999-03-19
FRIOR PILING DATE: 2000-09-20
FRIOR PILING DATE: 1999-03-19
                                                                                                                                                                                                                                      1226 AAAAACTCAAGCGGCTGGAAGGAGGAAGAGGTTCTCCAGAGTCGGAAC 1273
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Sequence 207, Application US/09925301

Sequence 207, Application US/09925301

Sequence 207, Application US/09925301

Settle No. US2002002308A1

SETTLE NO. US200200203304

FILE REFERENCE: Pal16

CURRENT APPLICATION NUMBER: US/09/925,301

CURRENT APPLICATION NUMBER: US/09/925,301

FRIOR PILING DATE: 2001-08-10

PRIOR FILING DATE: 2000-03-08

PRIOR FILING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 1694

SEQ ID NO 207

LENGTH: 1996
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Matches:
Conservative:
Mismatches:
Indels:
     Conservative:
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                         Mismatches:
Indels:
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ORGANISM: Homo sapiens
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US-10-131-410-64

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	Length:	Matches:	Conservative:	Mismatches:	Indels:	Gaps:
	3.63e-05	84.00	100.00\$	100.00%	100.00%	12
Alignment Scores:	Pred. No.:	Score:	Percent Similarity:	Best Local Similarity: 100	Query Match:	DB:

US-08-978-217-12 (1-16) x US-10-131-410-64 (1-2269)

Search completed: February 13, 2004, 01:46:08 Job time : 28.2461 secs

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Sequence 944, App Sequence 944, App Sequence 953, App Sequence 853, App Sequence 813, App Sequence 101, App Sequence 101, App Sequence 102, App Sequence 105, App Sequence 105, App Sequence 105, App Sequence 105, App Sequence 105, App Sequence 109194, Sequence 199194, Sequence 199194, Sequence 199197, Sequence 199196, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 199197, Sequence 697, App Sequence 697, App Sequence 697, App Sequence 697, App Sequence 191197, Sequence 697, App Sequence 191197, Sequence 697, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequence 10113, App Sequenc
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APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Gardon B.
APPLICANT: Mang, Aljun
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEC ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
                     ALIGNMENTS
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Patent No. US20020076414A1
GENERAL INFORMATION:
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Magher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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-D3=-bulished_Applications Na -QFWT=fastap - SUFFTX=rnpb - MIRMATCH=0.1
-LGOPEL=0 - LOOPEXT=0 - UNITS=bits - START=1 - END=1 - MATRIX=blosum6.2
-TRANS=human40.cdi - LIST=45 - DOCALIGN=20 - THR SCORE=pot - THR MXX=100
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-NOVEL6 - ICPU-3 - NO MMAP - LARGEQUERY - NEG SCORES=0 - WAIT - DSPBLOCK=100
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-FGAPOP=6 - FGAPEXT=7 - YGAPOP=10 - YGAPEXT=0.5 - DELOP=6 - DELEXT=7
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                          GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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Maximum Match 100%
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Match Length DB
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Sequence:
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412 CAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG 353
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412 CAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG 353
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                                                                                                                                      61 ProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHisProGlySer 80
                                                                                352 GAGAAGGATGGCATGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGG
                                                                                                                                                               41 GluLysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAspGlnGlySer
                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION;
APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Mangy Tongtong
APPLICANT: Wangy Trongtong
APPLICANT: Smith, Carole L.
APPLICANT: Smith, Carole L.
APPLICANT: Anny, Yasir A. W.
APPLICANT: Steiky, Yasir A. W.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Panger, Jonathan D.
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE REPREBUCE: 210121.471C14
CURRENT APPLICANT: VOICE NOTION NUMBER: US/10/025,380
CURRENT APPLICANT: Fanger: US/10/025,380
CURRENT PLING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SEO ID NOS: 1129
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Matches:
Conservative:
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                                                                                                                                                                                                                                                                                                                                   ; Sequence 944, Application US/10025380; Publication No. US20020182191A1; GENERAL INFORMATION:
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US-10-025-380-944
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Best Local Similarity:
Query Match:
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US-10-025-380-944/c
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Pred. No.:
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LENGTH: 563
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US-09-833-263-944/c

US-09-833-263-944/c

US-09-833-263-944/c

Sequence 944, Application US/09833263

Patent No. US2002011547A1

GENERAL INFORMATION:

APPLICANT: Anang, Aijun

APPLICANT: Clapper, Jonathan D.

APPLICANT: Stolk, John A.

APPLICANT: Macadler: Macadler: Macadlerin J.

TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

FILE REPRERENCE: 210.21.471C12

CURRENT APPLICATION NUMBER: US/09/833,263

CURRENT FILING DATE: 2001-04-10

NUMBER OF SEQ ID NOS: 1093

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      472 AATTGTGCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCC 413
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Matches:
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    ; SEQ ID NO 944
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: TYPE: DNA
; CRGANIEM: Homo sapiens
US-09-922-217-944
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CRGANISM: Homo sapien
US-09-833-263-944
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us-08-978-217-7.rnpb

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Sequence 853. Application US/10025380; Publication No. US20020182191A1; Publication No. US20020182191A1; APPLICANT: Xu, Jiangchun; APPLICANT: Lodges, Michael J. APPLICANT: Benson, Darin R. APPLICANT: Benson, Darin R. APPLICANT: Benson, Darin R. APPLICANT: Scolk, John A. APPLICANT: Wang, Tongtong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jiang, Yuqiu
Smith, Carole L.
King, Gordon E.
Wang, Ajun
Clapper, Jonathan D.
Skeiky, Yasir A. W.
Fanger, Gary R.
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   APPLICANT: Stolk, John A.
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CRGANISM: Homo sapien
US-09-833-263-853
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Pred. No.:
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Mismatches:
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CURRENT APPLICATION NUMBER: U8/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 853
LENGTH: 626
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Patent No. US20020110547A1
GENERAL INFORMATION:
APPLICANT: Wang, Aljun
APPLICANT: Clapper, Jonathan D.
                                                                                                                                                                                                APPLICANT: Xu, Jiangchun APPLICANT: Lodes, Michael J. APPLICANT: Becrist, Heather APPLICANT: Benson, Darin R. APPLICANT: Meagher, Madeleine Joy APPLICANT: Stolk, John A. APPLICANT: Wang, Tongtong
                                                                                                                           5-09-922-217-853/c
Sequence 853, Application US/09922217
Patent No. US20020076414A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jiang, Yudiu
APPLICANT: Smith, Carole Lynn
APPLICANT: King, Gordon E.
APPLICANT: Wang, Aijun
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US-09-922-217-853
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Best Local Similarity:
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APPLICANT: Meadler, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 21012.1-471C12
CURRENT APPLICATION NUMBER: U5/09/833,263
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOPTHARE: FastSEQ for Windows Version 3.0
LENGTH: 626
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CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Mindows Version 4.0
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APPLICANT: Machael MORRISEY
APPLICANT: Machael MORRISEY
APPLICANT: Peter OLANDT
APPLICANT: Peter VEIBY
APPLICANT: Peter VEIBY
APPLICANT: Peter VEIBY
APPLICANT: ROSERT C. BAST, Jr.
APPLICANT: RALE LU
APPLICANT: RALE LU
APPLICANT: RALE LU
APPLICANT: RALE CIAAT
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Nucleic Acid Molecules and Therapy of Ovarian Cancer
TITLE OF INVENTION: Nucleic Acid Molecules and Therapy of Ovarian Cancer
TITLE OF INVENTION: NUMBER: US/10/097,340
CURRENT APPLICATION NUMBER: 60/25,449
PRIOR APPLICATION NUMBER: 60/225,449
PRIOR PILING DATE: 2001-09-26
PRIOR PILING DATE: 2001-09-26
PRIOR PILING DATE: 2001-09-14
PRIOR PILING DATE: 2001-09-14
PRIOR PILING DATE: 2001-09-14
PRIOR PILING DATE: 2001-09-16
PRIOR PILING DATE: 2001-09-16
PRIOR PILING DATE: 2001-09-10
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PRIOR PILING DATE: 2001-09-10
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, Sequence 74, Application US/10097340
; Publication No. US20030087250A1
; GENERAL INFORMATION:
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APPLICANT: Marjula GANNAVRAPU
APPLICANT: Sebastian HOERSCH
APPLICANT: Steve G. KOVATS
APPLICANT: Rachel E. MEYERS
APPLICANT: Rachel E. MEYERS
APPLICANT: Michael MORRISEY
APPLICANT: Peter OLANDT
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                                      TYPE: DNA
ORGANISM: Homo sapiens
US-10-025-380-853
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Best Local Similarity:
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SEQ ID NO 853
LENGTH: 626
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RESULT 8

US-09-964-824A-101

Sequence 101, Application US/09964824A

Patent No. US20020102531A1

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: HOLTIGAN, Stephen

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signation of TITLE OF INVENTION: Set Sequence:

TITLE OF INVENTION: Cate Sequence:

TITLE OF INVENTION: Cate Sequence:

TITLE OF INVENTION: Cate Sequence:

TITLE OF INVENTION: Cate Sequence:

FILE PERRENCE: 689290-73

CURRENT APPLICATION NUMBER: US/60/236,033

PRIOR FILING DATE: 2000-09-28

PRIOR APPLICATION NUMBER: US/60/236,036

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR FILING DATE: 2000-09-28

NUMBER: US/60/236,028

NUMBER: OF SEQ ID NOS: S93

NUMBER: OF SEQ ID NOS: S93
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Matches:
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Mismatches:
Indels:
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PRIOR FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/323,580
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 1907
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SEQ ID NO 101
LENGTH: 1915
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ORGANISM: Homo sapiens
US-10-097-340-74
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Sequence 192, Application US/09967768A
Patent No. USZ0020150877A1
GENERAL INFORMATION:
APPELAL INFORMATION:
APPELAL INFORMATION:
TITLE OF INVENTION: Sets
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-72
CURRENT APPLICATION NUMBER: US/09/967,768A
CURRENT FILING DATE: 2001-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               429 AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCC 488
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APPLICANT: Workley, Joseph G.
APPLICANT: Workley, Joseph G.
APPLICANT: Gene Logic, Inc.
APPLICANT: Gene Logic, Inc.
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer;
FILE REPERENCE: 44921-5028-W0.
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT PILING DATE: 2000-106-14
PRIOR PILING DATE: 2000-06-14
PRIOR PILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PatentIn Ver: 2.1
SEQ ID NO 3420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Genbank Accession No. US20020142981A1 U73843
US-09-880-107-3420
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                                                                                                        US-09-880-107-3420
; Sequence 3420, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
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81 CysGlyAlaGly 84
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Best Local Similarity:
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US-09-967-768A-192
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TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
TITLE OF INVENTION: Sets
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR PELING DATE: 2000-09-28
PRIOR PELING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
SOFTWARE: Patentin version 3.0
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US-08-978-217-7 (1-84) x US-09-964-824A-101 (1-1915)
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Sequence 563, Application US/09964824A

; Patent No. US20020102531A1

; GENERAL INFORMATION:
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APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Carole Lynn
APPLICANT: Wang, Aliun
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210121.471C13
                                                                                                                                                                                                                                                                                                                                                                                             Conservative:
Mismatches:
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UNEART FILING DATE: 2001-08-03
NUMBER OS SEQ ID NOS: 1124
SOFTWARE: FASESEQ for Windows Version 4.0
PRIOR APPLICATION NUMBER: US/60/236,109
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR PILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,111
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 325
SOFTWARE: Patentin version 3.0
LENGTH: 1915
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Patent No. US20020076414A1
GENERAL INFORMATION:
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APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Maegher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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CORGANISM: Homo sapiens
US-09-922-217-1105
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; ORGANISM: Homo sapiens
US-09-967-768A-192
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Best Local Similarity:
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US-09-922-217-1105
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LENGTH: 1917
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431 AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCC 490
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE FILE REFERENCE: 210121-471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: PASESEQ FOR WINDOWS Version 4.0
SEQ ID NO 1105
LENGTH: 1917
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Matches:
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SENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Lodge, Michael J.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
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Skeiky, Yasir A. W.
Fanger, Gary R.
Vedvick Thomas S.
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
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                                        Percent Similarity:
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US-10-025-380-1105
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                                                               431 AATTGTGCCCTTGAGGAGCTGCGTCTGGGCTCTTGGGCCCTCTGGGGGACCAACTCCATGCC 490
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                                      1 AsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeuHisAla 20
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PAL33PI
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: US/US01/18569
PRIOR PILING DATE: 2001-06-07
PRIOR FILING DATE: 2000-06-07
PRIOR FILING DATE: 2000-06-07
PRIOR FILING DATE: 2000-06-07
PRIOR FILING DATE: 2010-06-07
SOFTWARE: Patentin Ver: 3.1
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Mismatches:
Indels:
US-08-978-217-7 (1-84) x US-10-025-380-1105 (1-1917)
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Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 756, Application US/10264049; Publication No. US20040005579A1; GENERAL INFORMATION:
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100.00%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
CORGANISM: Homo sapiens
US-10-264-049-756
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Best Local Similari
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US-10-264-049-756
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LENGTH: 1956
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570 GAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGC 629
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          510 CAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCCAGTTGGATCATTGAGCTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 AsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGlnLeuHisAla
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                                                                                                               APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REPREEMED: PA106
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1990-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 207
LENGTH: 1996
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Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length:
                                             ; Sequence 207, Application US/09925301
; Patent No. US20020052308A1
; GENERAL INFORMATION:
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CRGANISM: Homo sapiens
US-09-925-301-207
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Best Local Similarity:
RESULT 15
US-09-925-301-207
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DB:
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February 13, 2004, 01:27:22 ; Search time 88.8409 Seconds (without alignments) 10448.744 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUBLseq:*
3: /cgn2_6/ptodata/1/pubpna/DSO6_NEW_PUBLseq:*
4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
5: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
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13: /cgn2_6/ptodata/1/pubpna/USO9_NEW_PUB.seq:*
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18: /cgn2_6/ptodata/1/pubpna/USO0NEW_PUB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2449703 seqs, 1841816367 residues
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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252
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Sequence 944, App		Sequence 944, App	Sequence 853, App	Sequence 853, App	Sequence 853, App	Sequence 74, Appl	Sequence 101, App	Sequence 563, App	Seguence 3420, Ap	Sequence 192, App	Sequence 1105, Ap	Sequence 1105, Ap	Sequence 756, App	Sequence 207, App
	. QI	US-09-922-217-944	US-09-833-263-944	US-10-025-380-944	US-09-922-217-853	US-09-833-263-853	US-10-025-380-853	US-10-097-340-74	US-09-964-824A-101	US-09-964-824A-563	US-09-880-107-3420	US-09-967-768A-192	US-09-922-217-1105	US-10-025-380-1105	US-10-264-049-756	US-09-925-301-207
	Query Match Length DB	63 9	563 10	53 14	6 92	26 10	26 14	15	15 10	10	10	10	σ	14	12	σ
	Lengt									1915						
₩	Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
	Score	252	252	252	252	252	252	252	252	252	252	252	252	252	252	252
	Result No.	0 1	O O	ო ს	Ω 4	υ 2	o 0	7	Φ	O)	10	11	12	13	14	15

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Sequence 64, Appl Sequence 1740, App Sequence 2216, App Sequence 2216, App Sequence 280656, Sequence 280565, Sequence 280564, Sequence 280564, Sequence 280564, Sequence 280564, Sequence 15959, Sequence 15959, Sequence 1801, App Sequence 2266, Appl Sequence 3220, Appl Sequence 3220, App	DIAGNOSIS THEIR USE	gth 563; dels 0; Gaps 0; GGGACCAACTCCATGCC 60
US-10-131-410-64 US-09-967-701-4818 US-09-998-598-1740 US-09-998-598-1740 US-09-998-598-1740 US-10-101-510-356 US-10-027-632-280565 US-10-027-632-280565 US-10-027-632-280565 US-10-027-632-280565 US-10-027-632-280564 US-10-027-632-280564 US-10-027-632-280564 US-10-027-632-280564 US-10-027-632-159559 US-10-027-632-159559 US-10-027-632-159559 US-10-027-632-159559 US-10-027-632-15959 US-10-027-632-159559 US-10-027-632-159559 US-10-027-632-159559 US-10-027-632-159559 US-10-027-632-159559 US-09-923-876-4804 US-09-923-876-4804 US-09-954-456-55 US-09-954-456-1132 US-09-954-456-1132 US-09-954-456-1132 US-09-954-456-1132 US-09-954-456-1132 US-09-954-456-1132 US-09-954-456-1132 US-09-954-456-1132 US-09-954-456-1132 US-09-954-531-145 US-09-954-531-3220 US-09-954-531-3220	ALIGNMENTS 1922217 1 JOY 1 JOY 2 CANCER AND METHODS FOR (709/922,217) 03 Version 4.0	; Score 252; DB 9; Length; Pred. No. 2.1e-64; Indel 0; Mismatches 0; Indel CTGCGTCTTGGGGG
2552 100.0 2269 12649 98.8 12649 98.8 1265 100 12669 1269 1269 1269 1269 1269 1269 126	BSULT 1 S-09-922-217-944/C Sequence 944, Application US/09) Patent No. US20020076414A1 GENERAL INFORMATION: APPLICANT: Xu, Jiangchun APPLICANT: Lodes, Michael J. APPLICANT: Secrist, Heather APPLICANT: Secrist, Heather APPLICANT: Secrist, Madeleine APPLICANT: Secrist, Madeleine APPLICANT: Stolk, John A. APPLICANT: Stolk, John A. APPLICANT: Stolk, John A. APPLICANT: Stolk, John A. APPLICANT: Stolk, John A. APPLICANT: Stolk, John A. APPLICANT: Stolk, John A. APPLICANT: Gracle Lynn APPLICANT: Gracle APPLICANT: Gracle APPLICANT: Gracle APPLICANT: Gracle APPLICANT: Gracle APPLICANT: Wang, Ailun APPLICANT: Gracle APPLI	Match Local Similarity 100.08 ss 252; Conservative 1 AATTGTGCCTTGAGGAG
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Patent No. US20020110547A1

GENERAL INFORMATION:

APPLICANT: Wang, Aijun

APPLICANT: Clapper, John A.

APPLICANT: Clapper, John A.

TITLE OF INVENTION: CMPOUNDS FOR INMUNOTHERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS

TITLE OF INVENTION NUMBER: US/09/833,263

CURRENT APPLICATION NUMBER: US/09/833,263

CURRENT FILING DATE: 2001-04-10

NUMBER OF SEQ ID NOS: 1093

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 944

LENGHAL: 563
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                            412 CAGCTGCGAGACCTCACTTCCAGTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG 353
                                                                                                                                                                                                      292 CCCTTTGCCCAGGAGCTGCTGGACGGTCAGCAGCCAGCAGCCCCTACCACCCGGCAGC 233
61 CAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG 120
                                                                                        121 GAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGG
                                                                                                                                352 dadaaddarddcarddccrrccaddaddcccradacccadddccrrrrgaccaddcaddcad
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Best Local Similarity 100.0
Matches 252, Conservative
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                                                                                                                                                                                                                                                                                                          232 TGTGGCGCAGGA 221
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; ORGANISM: Homo sapien
US-09-833-263-944
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US-10-025-380-944/c
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Sequence 944, Application US/10025380; Publication No. US20020182191A1; GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.

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412 CAGCTGCGAGACCTCACTTCCAGCTCTTGAGCTCAGTTGGATCATTGAGCTGCTG 353
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT FALLING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
                                                                                                                                                                                                                                                   APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COMPOUNDS
FILE REPERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILIG DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 944
LENGTH: 563
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Patent No. US20020076414A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
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Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
Jiang, Yuqiu
Smith, Carole Lynn
King, Gordon E.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
                                                                    Jiang, Yuqiu
Smith, Carole L.
King, Gordon E.
Wang, Aljun
Clapper, Jonathan D.
Skeiky, Yasir A. W.
Fanger, Gary R.
Vedvick Thomas S.
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US-10-025-380-944
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471 AATRGRGCCCTTGAGGAGCTGCGTCTTGGGGCCTCTGGGGGGACCAACTCCATGCC 412
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APPLICANT: Clapper, John A.
APPLICANT: Stalk, John A.
APPLICANT: Stalk, John A.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C12
CURRENT APPLICANTION NUMBER: US/09/833,263
CURRENT APPLICANTION NUMBER: US/09/833,263
SOFTWARE: FastSEQ for Windows Version 3.0
IENGTH: 626
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SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 853 LENGTH: 626
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; Patent No. US20020110547A1
; GENERAL INFORMATION:
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                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
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US-09-833-263-853
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.41/214
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FASISED for Windows Version 4.0
SEQ ID NO 853
LENGTH: 626
                                                                                                                                                                                                                                                         APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Becrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Program
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Sinth, Carole L.
APPLICANT: Smith, Carole L.
APPLICANT: Smith, Carole L.
APPLICANT: Smith, Carole L.
APPLICANT: Smith, Gordon B.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Fanger, Jonathan D.
APPLICANT: Fanger, Jonathan B.
APPLICANT: Fanger, Jonathan B.
APPLICANT: Fanger, Jonathan B.
APPLICANT: Fanger, Jonathan B.
APPLICANT: Carter, Darrick
                                                                                                                                                                                    Sequence 853, Application US/10025380 Publication No. US20020182191A1 GENERAL INFORMATION:
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241 TGTGGCGCAGGA 252
                                                   231 TGTGGCGCAGGA 220
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Sequence 563, Application US/09964824A

Sequence 563, Application US/09964824A

Parent No. US20020102531A1

GENERAL INFORMATION:

TITLE OF INVENTION: Gancer Gene Determination and Therapeutic Screening Using Signation TITLE OF INVENTION: Sets

FILE REPRENCE: 689250-73

CURRENT APPLICATION NUMBER: US/09/964,824A

CURRENT APPLICATION NUMBER: US/60/236,033

PRIOR FILING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28

PRIOR PELING DATE: 2000-09-28

PRIOR PILING DATE: 2000-09-28
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Best Local Similarity 100.0%; Pred. No. 2.4e-64;
Matches 252; 'Conservative 0; Mismatches 0; Indels 0;
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FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
SOFTWARE: PatentIn version 3.0
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ORGANISM: Homo sapiens
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COCGANISM: Homo sapiens
US-09-964-824A-563
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    Sequence 101, Application US/0964824A
    Patent No. US2020102531A1
    GENERAL INFORMATION:
    APPLICANT: Horrigan, Stephen
    TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatulation Final Properties

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TITLE OF INVENTION: Nucleic Acid Molecules and Proceins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
FILE REFERENCE: MRI-030
CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR PILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR PILING DATE: 2001-03-14
PRIOR PILING DATE: 2001-03-16
PRIOR PILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/324,967
PRIOR PILING DATE: 2001-03-16
PRIOR PILING DATE: 2001-09-26
PRIOR PILING DATE: 2001-09-26
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PRIOR PILING DATE: 2001-09-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 363
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                       Gordon B. MILLS
Robert C. BAST, Jr.
          Shubhangi KAMATKAR
                                                                                                                                                                                                                                                                                                                                       Rosemarie SCHMANDT
                                                                       Rachel E. MEYERS
Michael MORRISEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   241 TGTGGCGCAGGA 252
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                                          KOVATS
                                                                                                                                          Peter OLANDT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-097-340-74
                                                                                                                                                                                                           Peter VEIBY
                                                                                                                                                                                                                                                                                                                                                                           Xumei ZHAO
Karen GLATT
                                                                                                                                                                                                                                                                                                           Karen LU
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US-09-964-824A-101
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LENGTH: 1907
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APPLICANT:
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Sequence 192, Application US/09967768A
Patent No. US2020150877A1
GENERAL INFORMATION:
APPLICANT: Augustus, Meena
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Secrist, Heather
Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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US-09-967-768A-192
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APPLICANT:
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    429 AATIGIGCCCTIGAGGAGCIGCGICTGIGGICTITGGGCCTCTGGGGGACCAACTCCAIGCC 488
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 3420, Application US/09880107
; Patent No. US2020142981A1
; GENERAL INFORMATION:
    APPLICANT: Horne, Darci T.
    APPLICANT: Horne, Darci T.
    APPLICANT: Gene Logic, Inc.
    TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer; TILE OF INVENTION: Gene Expression Profiles in Liver Cancer; TILE OF INVENTION: Gene Expression Profiles in Liver Cancer; TILE OF INVENTION: Gold Expression NUMBER: US/09/880,107
; CURRENT APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR PILING DATE: 2000-06-14
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 3950
; NUMBER OF SEQ ID NOS: 3950
; SOFTWARE PATENTIN VET. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0%; Score 252; DB 10; Length 1915;
Best Local Similarity 100.0%; Pred. No. 2.4e-64;
Matches 252; Conservative 0; Mismatches 0; Indels 0;
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US-09-880-107-3420
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US-09-967-768A-192
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LENGTH: 1915
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ATTLE OF INVENTION: America TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signaturity of INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF INVENTION: Sets TITLE OF 
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Patent No. US20020076414A1
GENERAL INFORMATION:
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Query Match
Best Local Similarity 100.0
Matches 252; Conservative
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100.0%; Pred. No. 2.4e-64;
vative 0; Mismatches 0; Indels 0;
                                                                                           Query Match 100.0%; Score 252; DB 9; Length 1917; Best Local Similarity 100.0%; Pred. No. 2.4e-64; Matches 252; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 1129
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 1105
LENGTH: 1917
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Publication No. US20020182191A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Macher, Madeleine Joy
APPLICANT: Stolk, John A.
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Skeiky, Yasir A. W.
Fanger, Gary R.
Vedvick Thomas S.
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Smith, Carole L.
King, Gordon E.
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Matches 252; Conservative
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US-10-025-380-1105
                TYPE: DNA
ORGANISM: Homo sapiens
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Best Local Similarity
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LENGTH: 1917
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 756, Application US/10264049
; Publication No. US2004005579A1
; GENERAL INFORMATION:
APPLICANT: Birse et al.
; IILLE OF INVENTION: Mucleic Acids, Proteins, and Antibodies
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US-09-925-301-207
Squarece 207, Application US/09925301
Fatent No. US20020052308A1
GENERAL INFORMATION:
FAPLICANT: ROSEN et al.
FILE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA106
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: PA137P1
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/18569
PRIOR FILING DATE: 2001-06-07
PRIOR FILING DATE: 2000-06-07
PRIOR FILING DATE: 2000-06-07
NUMBER: OF SEQ ID NOS: 4360
SEQ ID NO 756
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450 AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCC 509
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100.0%; Score 252; DB 9; Length 1996;

Best Local Similarity 100.0%; Pred. No. 2.4e-64;

Matches 252; Conservative 0; Mismatches 0; Indels 0
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: Pacentin Ver. 2.0
ERQ ID NO 207
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CAGANISM: Homo sapiens
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Search completed: February 13, 2004, 11:48:19 Job time : 89.8409 secs

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APPLICANT: Gordon B. MILLS
APPLICANT: Robert C. BAST, Jr.
APPLICANT: Karen LU
APPLICANT: Karen LU
APPLICANT: Xumei ZHAO
APPLICANT: Xumei ZHAO
APPLICANT: Karen GLATT
TILLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer FILE REFERENCE: MRI-030
   Sequence 74, Appl Sequence 101, App Sequence 1105, App Sequence 1105, App Sequence 1105, App Sequence 1105, App Sequence 207, App Sequence 207, App Sequence 283, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 282, App Sequence 309, App Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence 44, App1 Sequence
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Sequence 44, Appl
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Sequence 11873, A
15 US-10-097-340-74
10 US-09-964-824A-101
10 US-09-964-824A-101
10 US-09-964-824A-192
11 US-10-26-922-217-105
12 US-10-26-922-217-105
13 US-09-922-217-105
14 US-10-025-380-1105
15 US-09-922-217-1944
16 US-09-922-217-944
17 US-09-922-380-944
18 US-09-922-380-944
19 US-09-922-380-944
19 US-09-922-380-944
19 US-09-922-217-944
19 US-09-922-2180-944
19 US-09-922-282
11 US-09-922-282
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22 US-09-922-798-1601
23 US-09-922-798-1601
24 US-09-113-444
25 US-09-923-880-44
26 US-09-921-828-44
27 US-09-113-828-14
28 US-10-010-940-44
28 US-10-010-940-44
29 US-09-926-328-126
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; Sequence 74, Application US/10097340
; Publication No. US20030087250A1
; GENERAL INFORMATION;
APPLICANT: John MONAHAN
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Sebastian HOERSCH
Shubhangi KAMATKAR
Steve G. KOVATS
Rachel E. MEYERS
Michael MORRISEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peter VEIBY
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-MODEL=frame+ p2n.model -DEV=x1h
-MODEL=frame+ p2n.model -DEV=x1h
-Q=/cgn2_1/USFTO_spool/US08978217/runat_10022004_133827_20514/app_query.fasta_1.1500
-Q=/cgn2_1/USFTO_spool/US08978217/runat_10022004_133827_20514/app_query.fasta_1.1500
-DB=Published Applications NA -QFMT=fastap -SUFFTX=rnpb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -BND=1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR SOXE=pstc -THR MAXEN-2
-TTRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR SOXES=pstc -THR MAXEN-2
-THR MIN=0 -ALIGN=15 -MODE=LOCAL -UOTFPMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXIEN=200000000 -USER=US08978217 @CGN 1 1 690 @runat 10022004 133827_20514
-NCPUG= -ICCPU=3 -NO PMAP -LARGEQUERS -NEG SOCRES=0 -WAIT -DSBLGCK=100
-LONGLOG -DEV TIMEOÜT=120 -WARN TIMEOUT=3 -THRABAD=1 -XGAPDP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                     February 12, 2004, 21:34:51 ; Search time 585.395 Seconds (without alignments) 2334.540 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                          US-08-978-217-2
1980
1 MAATCEISNIFSNYFSAMYS.......YKFGKNSSGWKEEEVLQSRN 371
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/PCT_NEW PUB.seq:*
3: /cgn2_6/ptodata/1/pubpna/US06_DNB/W PUB.seq:*
4: /cgn2_6/ptodata/1/pubpna/US06_DNB/COMB.seq:*
5: /cgn2_6/ptodata/1/pubpna/US06_DNB/COMB.seq:*
6: /cgn2_6/ptodata/1/pubpna/US07_NEW PUB.seq:*
7: /cgn2_6/ptodata/1/pubpna/US08_DUBCOMB.seq:*
8: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
9: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
10: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
11: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
12: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
13: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
14: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
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16: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
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                                       GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                        OM protein - nucleic search, using frame_plus_p2n model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2449703 seqs, 1841816367 residues
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Maximum Match 100%
Listing first 45 summaries
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Xgapop 10.0, Xgapext
Ygapop 10.0, Ygapext
Fgapop 6.0, Fgapext
Delop 6.0, Delext
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Maximum DB seq length: 2000000000
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1056 AACAGCAACATGACCTACGAGAAGCTGAGGAGCCGGGCCATGAGGTACTACTACAAACGGGAG 1115
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                                                 201 ThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySerAspValAspLeuAsp 220
                                                                                                                                                                                                                                                                                                                                                                                                                   261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         281 AspileLeulleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    936 GACATCCTCATCCACCCGGAGCTCAACGAGGCCTCATGAAGTGGGAGAATCGGCATGAA 995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  301 GlyvalPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLys 320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        321 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrLysArgGlu 340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              341 IleLeuGluArgValAspGlyArgArgLeuValTyrLysPheGlyLysAsnSerSerGly 360
                                                                                           696 ACTGGTGCTTCTCCGAGCTCCCACTCCTCAGACTCCGGTGGAAGTGACGTGGACCTCGACT
                                                                                                                                                                                                                        756 CCCACTGATGGCAGCTCTTCCCCAGCGATGGTTTTCGTGACTGCAAGAAGAAGGGCGGATCCC
                                                                                                                                                                                                                                                                                             LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys
                                                                                                                                                                                                                                                                                                                                           816 AAGCACGGGAAGCGGAAACGAGGCCGGCCCCGAAAAGCTGAGCAAAAAGGTACTGGGACTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                    221 ProThrAspGlyLysLeuPheProSerAspGlyPheArgAspCysLysLysGlyAspPro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         161 GinglySerProPheAlaGInGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHis 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    635
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         636 CCCGGCAGCTGTGGCGCAGGAGCCCCCTCCCCTGGCAGCTCTGACGTCTCCACCGCAGGG 695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          516 GAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGAC 575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96 ATGGCTGCAACCTGTGAGATTAGCAACATTTTTAGCAACTACTTCAGTGCGATGTACAGC 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         156 TCGGAGGACTCCACCCTGGCCTCTGTTCCCCCTGCTGCCACCTTTGGGGCCCGATGACTTG 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       81 GlurysAspirysTyrAspAlaSerAlaIleAspPheSerArgCysAspMetAspGlyAla 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  101 ThrLeuCysAsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGln 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    396 ACCCTCTGCAATTGTGCCCTTGAGGAGCTGCGTCTGGGTCTTTGGGCCTCTGGGGACCAA 455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   216 GTACTGACCCTGAGCAACCCCCAGATGTCATTGGAGGGTACAGAGAAAAAGGCCAGCTGGTTG 275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         276 GGGGAACAGCCCCAGTTCTGGTCGAAGACGCAGGTTCTGGACTGGATCAGCTACCAAGTG 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        41 ValLeuThrLeuSerAsnProGlnMetSerLeuGluGlyThrGluLysAlaSerTrpLeu 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       456 CICCAIGCCCAGCIGCGAGACCICACINCCAGCICINCIGAIGAGCICAGIIGGAICAIT
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Matches:
Conservative:
Mismatches:
Indels:
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CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: 00/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-03-14
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Pred. No.:
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Sequence 101, Application US/09964824A

Sequence 101, Application US/09964824A

Between No. US20020102531A1

GENERAL INFORMATION:

APPLICANT: HOTIGAN: Stephen

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat:

TITLE OF INVENTION: Set 8

FILE REFERENCE: 689200-73

CURRENT APPLICATION NUMBER: US/60/236,033

FRIOR PELLING DATE: 2000-09-28

PRIOR FILING DATE: 2000-09-28

PRIOR APPLICATION NUMBER: US/60/236,032

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR APPLICATION NUMBER: US/60/236,028

PRIOR PILING DATE: 2000-09-28

NUMBER OF SEQ ID NOS: 583

SOFTWARE: Patentin version 3.0

SEQ ID NO 101

LENGTH: 1915
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Matches:
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Db 1200 TGGAAGGAGGAAGGTTCTCCAAGG	KESULT 3 US-904-824A-563 ; Sequence 563, Application US/09964824A ; Patent No. US20020102531A1	; APPLICANT: Horrigan, Stephen ; TITLE OF INVENTION: Cancer Gene Determ ; TITLE OF INVENTION: Sets	; FILE REFERENCE: 09290'3; CURRENT APPLICATION NUMBER: US/09/964, CURRENT FILING DATE: 2001-09-27; PRIOR APPLICATION NUMBER: US/60/236,03	; PRIOR FILING DATE: 2000-09-28 ; PRIOR APPLICATION NUMBER: US/60/236,03 ; PRIOR FILING DATE: 2000-09-28 ; PRIOR APPLICATION NUMBER: US/60/236,02	FALOR FILING DALE: SOUTH STATES SOUTHWER OF SEQ ID NOS: 583 ; SOFTWARE: Patentin version 3.0 ; SEQ ID NO 563	TYPE: DNA ORGANISM: H S-09-964-824A-	8,69e-212 1980.00	Percent Similarity: 100.00* Co Best Local Similarity: 100.00* Mi Query Match: 100.00* In DB: 69	US-08-978-217-2 (1-371) x US-09-964-824A- Qy	Db 120 ATGGCTGCAACCTGTGAGATTAGCAAC Qy 21 SerGluAspSerThrleualaSerVal	180	Db 240 GTACTGACCCTGAGGAACCCCAGATG	Qy 61 GlyGluGlnProGlnPheTrpSerLys	Cy 81 GluLysAsnLysTyrAspAlaSerAla	Db 360 dadahdhachadhachcdchadcdccc	Db 420 ACCCTCIGCAALTIGTGCCCTTGAGGAG	121	Db 480 CTCCATGCCCAGCTGCGAGACCTCACT Oy 141 GluLeuLeuGluLysAspGlyMetAla	Db 540 GAGTGCTGGAGAAGGATGGCATGGCTGGCTGGAGGATGGCTGGAGAGGAAGGA	009	
	21 SerGluAspSerThrLeuAlaSerValProProAlaAlaThrPheGlyAlaAspAspLeu 40	41 ValleuThrleuSerAsnProGlnMetSerleuGluGlyThrGluLysAlaSerTrpleu 60 	61 GlygluGlnProGlnPheTrpSerLysThrGlnValLeuAspTrpIleSerTyrGlnVal 80 	81 GluiysAsniysTyraspalaSeralaileaspPheSerargCysAspMetaspGlyala 100 	101 ThrieucysAsnCysAlaieuGlucluleuArgleuValPheGlyBroLeuGlyAspGln 120	121 LeuHisAlaGlnLeuArgAspLeuThrSerSerSerSerAspGluLeuSerTrp1le11e 140	141 GluLeuLeuGluLysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAsp 160 	161 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHis 180 	181 ProdlySerCysGlyAladlyAlaBroSerProdlySerSerAspValSerThrAladly 200	201 ThrdlyAlaSerArgSerSerHisSerSerAspSerGlyGlySerAspValAspLeuAsp 220	ProThrAspGlyLysLeuPheProSerAspGlyPheArgAspCysLysLysGlyAspPro	780 CCCACTGAIGGCAAGCTCTTCCCCAGCGAIGGTTTTCGIGACTGCAAGAAGGGGGGATCCC 839 241 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys 260	840 AAGCAGGAAGCGGAAACGAGGCCGCCCCGAAAGCTGAGCAAAGATACTGGGACTGT 899 261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPheIleArg 280		281 AspIleLeulleHisProGluLeuAsnGluGlyLeuWetLygTrpGluAsnArgHisGlu 300	301 GlyValPheLygPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLygLygFyg 320	GGGTCTTCAAGTTCCTGCGCTCCGAGGCTGTGGCCCCAACTATGGGGCCAAAAAAAA	321 AbnSerabnMetThrTyrGluLysLeuSerArgalaMetArgTyrTyrTyrLygArgGlu 340 	341 IleLeuGluArgValAspGlyArgArgLeuValTyrIvsPheGlyLysAsnSerSerGly 360 	TrpLysGluGluGluValLeuGlnSerArgAsn 371	
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DB: US-08-978-217-2 (1-371) x US-09-880-107-3420 (1-1915) OY 1 MetalaalaThrCysGluIleSerAsnIlePheSerAsnTyrPheSerAlaMetTyrSer 20	TGCCACCTTTGGGGCCGATC TGCCACTTTGGGGCCGATC UG1yThrGluLysAlaSer1	Oy 61 GlyGluGlnProGlnPheTrpSerLysThrGlnValLeuAspTrpIleSerTyrGlnVal 80	Oy 81 GlulyshsniysTyraspalaSeralaileAspPheSerargCysAspMetAspGlyala 100	Oy 101 ThrLeucysAsnCysAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGln 120	Oy 121 LeuHisAlaGlnLeuArgAspLeuThrSerSerSerSerSerAspGluLeuSerTrpIleIle 140	Qy 141 GluleuleuGluLysAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAsp 160 Db 540 GAGCTGCTGCAGAAGGATGGCCATCCAGGAGGCCCTAGACCCAGGGCCCTTTGAC 599	Oy 161 GlnGlySerProPheAlaGlnGluLeuLeuAspaspGlyGlnGlnAlaSerProTyrHis 180	Oy 181 ProGlySerCysGlyAlaGlyAlaProSerProGlySerSerAspValSerThrAlaGly 200- 	Qy 201 ThrGlyAlaSerArgSerSerHisSerSerHisSerSerHisS 220 1	Qy 221 ProThrAspGlyLysLeuPheBroSeraspGlyPheArgAspCysLysLysLysGlyAspPro 240 1	CY 241 LYSHISG1YLYSARGLYSARGG1YARGPROARGLYSCHUSERLYSG1uTYTTTDASDCYS 260 B40 AAGCACGGGAAAGCGGAAACGGAGCCGCGCCCCGAAAGCTGAGAAAAGAGACTGT 899	Oy 261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPhelleArg 280	Cy 281 AspileLeuileHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu 300	Oy 301 GlyValPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLys 320	321 AsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArgTyrTyrTyrLysArgGlu 3	
aG1y 200 	CCC 839 CCZ 8260 	Arg 280 GG 959	Glu 300 	Lys 320 AAG 1079	ArgGlu 340 CGGGAG 1139	rGly 360 CGGC 1199									-	<u> </u>
7 7 3 7 7	CCCACTGATGGCAAGCTCTTCCCCAGCGATGGTTTTCGTGACTGCAAAAGGGGGATCCC LyshisGlyusAtgLysktgGlyatgProArgProArgusEusSetlysGlutTrpAspCys LyshisGlyly	LeuglugiyiysiysSeriyysHisAlaProArgglyThrHisLeuTrpgluPheIleArg 	ASPIJELEUIIEHISPROGIULEUASMGIUGIYLEUMETLYSTRPGIUASMARGHISGIU 	GlyvalPhelysPheLeuargSerGlualavalAlaGinLeuTrpGlyGlnLysLysEvsEvs 	AsnSerasnwetThrTyrGlulysLeuserargalametargTyrTyrTyrLysArgC 	- - - - - - - - - - - -	nSerargasn 371 GaGTCGGAAC 1232	107	<pre>/ PAGENT NO. USZUUZUL42981A1 / GENERAL INFORMATION: / APPLICANT: Horne, Darci T. / APPLICANT: Vockley, Joseph G.</pre>	n Profiles in Liver Cancer	880,107 1,379	7,054		iton No. US20020142981A1 U73843	Length: 1915 Matches: 371 Conservative: 0	
ProGlySerCysGlyAlaGlyAla 	CCACTGATGGCAAGCTCTTCCCC LyshisGlyLysargGly AGCACGGAAGCGAAACGGAAACGAGG	LeuGludlyLysLysSerLysHis 	Aspileteuilehisprogluleu 	GlyvalPheLysPheLeuArgSer 	AsnSerasmMetThrTyrGluLys 	IleLeuGluArgValAspGlyArc 	TrpLysGluGluGluValLeuGlnSerArgAsn 	3420 0, Application US/098801	SZUUZU1429BIAI RMATION: Horne, Darci T. Vockley, Joseph G.	Schert, Uwe Gene Logic, Inc. VENTION: Gene Expression NCE: 44921-5028-WO	LICATION NUMBER: US/09/8 ING DATE: 2001-06-14 CATION NUMBER: US 60/211 G DATE: 2000-06-14	CATION NUMBER: US 60/23° G DATE: 2000-10-02 EQ ID NOS: 3950 atentIn Ver. 2.1	O ID NO 3420 LENGTH: 1915 TYPE: DNA ORGANISM: Homo gapiens	FEATURE: OTHER INFORMATION: Genbank Accession No. 19-880-107-3420	es: 8.69e-212 1980.00 rity: 100.00\$	
Oy 181 b 660 C Oy 201 T	Db 780 (OX 241)	QY 261 1 Db 900 0	Qy 281 J	Oy 301 O	Oy 321 Db 1080	Oy 341 Db 1140	Oy 361 Db 1200	RESULT 4 US-09-880-107- ; Sequence 342	GENERAL INFC APPLICANT: APPLICANT:	APPLICANT: APPLICANT: TITLE OF IN; FILE REFERE	CURRENT APP CURRENT FIL: PRIOR APPLI	; PRIOR APPLI ; PRIOR FILIN ; NUMBER OF S ; SOFTWARE: P	; SEQ ID NO 3420 ; LENGTH: 1915 ; TYPE: DNA ; ORGANISM: HO	S-0	Alignment Scores Pred. No.: Score: Percent Similari	Best Local Similarity: Query Match:

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960 GACATCCTCATCCACCCGGAGCTCAACGAGGGCCTCATGAAGTGGGAAATCGGCATGAA 1019
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                              161 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHis 180
                                                                                                                                                                                                                                                                    720 ACTGGTGCTTCTCGGAGCTCCCACTCGTCAGACTCGGGTGGAAGTGACGTGGACCTGGAT 779
                                                             241 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCys
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       281 AspileLeulleHisProGluLeuAsnGluGlyLeuMetLysTrpGluAsnArgHisGlu
                                                                                                                       181 ProGlySerCysGlyAlaGlyAlaProSerProGlySerSerAspValSerThrAlaGly
                                                                                                                                                                   201 ThrGlyAlaSerArgSerSerHisSerSerAspSerGlyGlySerAspValAspLeuAsp
                                                                                                                                                                                                                                                                                                                    221 ProThrAspGlyLysLeuPheProSerAspGlyPheArgAspCysLysLysGlyAspPro
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| Sequence 1105, Application US/09922217
| Patent NO. USZ0020076414A1
| GENERAL INFORMATION|
| APPLICANT: No. Jiangchun
| APPLICANT: Lodes, Michael J.
| APPLICANT: Lodes, Michael J.
| APPLICANT: Benson, Darin R.
| APPLICANT: Benson, Darin R.
| APPLICANT: Broik, John A.
| APPLICANT: Stolk, John A.
| APPLICANT: Wang, Tongtong
| APPLICANT: Wang, Yuqiu
| APPLICANT: Wing, Yuqiu
| APPLICANT: Smith, Carole Lynn
| APPLICANT: Smith, Carole Lynn
| APPLICANT: Smith, Carole Lynn
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| APPLICANT: Smith, Carole Lynn
| APPLICANT: Smith, Carole Lynn
| APPLICANT: Tilly Gordon E.
| APPLICANT: Wang, Aijun
| APPLICANT: Wang, Aijun
| APPLICANT: TILLY OF INVENTION: COFCOLON CANCER AND DIAGNOSIS
| TITLE OF INVENTION: COFCOLON CANCER AND METHODS FOR THEIR USF
| TILLE REPERENCE: 210121.471C13
| CURRENT FILING DATE: 2001-08-03
| NUMBER OF SEQ ID NOS: 1124
| SEQ ID NO 1105
| LEMCTH: LEMCTH: 1917
| TUBERTH: NUMBER OF SEQ IC Windows Version 4.0
| EMROTH: NUMBER OF SEQ IC NOS: 1124
| TUBERTH: REALESQ for Windows Version A.0
| TUBERTH: NUMBER OF SEQ ID NOS: 1124
| TUBERTH: REALESQ for Windows Version A.0
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ORGANISM: Homo sapiens
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TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
TITLE OF INVENTION: Sets
FILE REPERBACE: 689290-72
CURRENT APPLICATION NUMBER: US/09/967,768A
CURRENT APPLICATION NUMBER: US/60/236,109
PRIOR PILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,034
PRIOR FILING DATE: 2000-09-28
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RIOR FILING DATE: 2000-09-28
RUMBER OF SEQ ID NOS: 325
SOFTWARE: PatentIn version 3.0
SEQ ID NO 192
LENGTH: 1915
1140 AICCTGGAACGGGTGGATGGCCGGCGACTCGTCTACAAGTTTGGCAAAAACTCAAGCGGC 1199
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                                                                                                                                                                                                           Sequence 192, Application US/09967768A Patent No. US20020150877A1 GENERAL INFORMATION:
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CORGANISM: Homo sapiens
US-09-967-768A-192
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	SULT 7 SOURT 7 -10-025-380- Sequence 110 Publication Publication	APPLICANT: Xu, Jiangchun APPLICANT: Lodes, Michael J. APPLICANT: Lodes, Michael J. APPLICANT: Berist, Heather APPLICANT: Benson, Darin R. APPLICANT: Meagher, Madeleine Joy APPLICANT: Stolk, John A. APPLICANT: APPLICANT: Tongtong APPLICANT: Jiang, Yuqiu	APPLICANT: King, Gordon E. APPLICANT: King, Gordon E. APPLICANT: Wang, Aljun APPLICANT: Clapper, Jonathan D. APPLICANT: Skeiky, Yasir A. W. APPLICANT: Panger, Gary R. W. APPLICANT: Panger, Gary R. S. APPLICANT: Vedvick Thomas S.	HAPPLICANT: CARLER, DARTICK TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE TITLE REFERENCE: 210121,471C14 CURRENT APPLICATION NUMBER: US/10/025,380 CURRENT FILING DATE: 2001-12-19 NUMBER OF SEQ ID NOS: 1129 SOFTWARE: FASTSEQ for Windows Version 4.0	ID NO 1105 PE: 1917 PE: DNA PANISM: Homo sapiens 125-380-1105 PE: Cores: 8.71e-212 Length:	1980.00 Similarity: 100.00\$ h: 100.00\$ h: 14	Oy 1 MetalahlaThrCysGluIleSerAsnIlePheSerAsnTyrPheSerAlaMetTyrSer 20
ignment Scor ed. No.: cet. Simila rcent Simila st Local Sim ery Match:	Oy 1 MetAlahlaThrCysGluIleSerAsnIlePheSerAsnTyrPheSerAlaMetTyrSer 20	41 ValleuThrLeuSerAsnProGlnMetSerLeuGluGlyThrGluLysAlaSerTrpLeu	Oy 81 GluLyBAbnLysfyrAspAlaSerAlaileAspPheSerArgCysAspMetAspGlyAla 100 Db 362 GAGAAGAACTACGACGCATACATTCTTCTACATGATGATGACGCC 421 CY 101 ThrLeuCyBAbnCyBAlaLeuGluGluLeuArgLeuValPheGlyProLeuGlyAspGln 120 Db 422 ACCTCTGCAATTGTGCCTTGAGGAGCTGCGTCTGGGCTCTTGGGGCTCTGGGGACCAA 481	Oy 121 LeuHisAlaGInLeuArgAspLeuThrSerSerSerAspGluLeuSerTrpIleIle 140 Db 482 CTCCATGCCAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGATCATT 541 Qy 141 GluLeuLeuGluLySAspGlyMetAlaPheGlnGluAlaLeuAspProGlyProPheAsp 160 Db 542 GAGCTGCTGGAGAAGGATGGCATGGCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGAC 601	CY 161 GlnGlySerProPheAlaGlnGluLeuLeuAspAspGlyGlnGlnAlaSerProTyrHis 180 Db 602 CAGGGCACCTTTTGCCCAGGAGCTGCTGGACGGTCAGCAAGCCAGCC	201 ThrGlyAlaGerArgSerSerHisSerSerAspSerGlyGlyGsrAspValAspLeuAsp	241 LysHisGlyLysArglyArgGlyArgProArglysLeuSerLysGluTyrTrpAspCvg 241 LysHisGlyLysArgLysArgGlyArgProArgLysLeuSerLysGluTyrTrpAspCvg 842 AAGCAGGAAGCGAAACGGGCCCCGGAAACTGGCAAAGAGTACTGGACTGT 261 LeuGluGlyLysLysSerLysHisAlaProArgGlyThrHisLeuTrpGluPherlaArg 902 CTCGAGGCAAGAAGAGCACGCCCCAGAGGCACCCCTGTGGAGTTCATCCGG 281 ASpIleLeuIleHisProGluLeuAsmGluGlyLeuMetLysTrpGluAsnArgHisGlu 962 GACATCCTCATCCACGCGGAAGAGCACCCCCTGTGGAAGTTCATCGG 301 GlyValPheLysPheLeuArgSerGluAlaValAlaGlnLeuTrpGlyGlnLysLysLys

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Publication No. US20040005579A1
GENERAL INFORMATION:
TILLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REPRENCE: PA133P1
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/18569
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                      60/209,467
PRIOR FILING DATE: 2001-06-07
PRIOR APPLICATION NUMBER: US 60,
NUMBER OF SEQ ID NOS: 4360
SOFTWARE: Patentin Ver. 3.1
SEQ ID NO 756
LENGTH: 1956
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Publication No. US20030235915A1
GENERAL INFORMATION
APPLICANT: SPECHT, THOMAS
APPLICANT: HINZMANN, BERND
APPLICANT: HINZMANN, BERND
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: POSSWTHAL, ANDRE
APPLICANT: ROSSWTHAL, ANDRE
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMAN SECUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMAN SECUENCES FROM TISSUE OF BREAST
FILE REFERENCE: SCH-1763
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Patent No. US20020052308A1
GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/131,410
CURRENT FILING DATE: 2002-04-25
FRIOR APPLICATION NUMBER: 09/646,673
FRIOR FILING DATE: 2000-09-20
FRIOR APPLICATION WUMBER: PCT/DE99/00908
FRIOR FILING DATE: 1996-03-19
NUMBER OF SEQ ID NOS: 202
SOFTWARE: Petentin Ver: 2.1
LENGTH: 2269
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ORGANISM: Homo sapiens
US-10-131-410-64
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                                                                                 315 TrpGlyGlnLysLysLysAsnSerAsnMetThrTyrGluLysLeuSerArgAlaMetArg
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APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrifst, Heather
APPLICANT: Berist, Heather
APPLICANT: Berist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Sinch, Carole Lynn
APPLICANT: Sinth, Carole Lynn
APPLICANT: Sinth, Carole Lynn
APPLICANT: Mang, Aijun
APPLICANT: Mang, Aijun
APPLICANT: The OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SEQ ID NO 953
LENTH: 626
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LENTH: 626
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Score: 1127.00
Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 56.92%
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CORGANISM: Homo sapiens
US-09-922-217-853
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133 SerAspGluLeuSerTrpIleIleGluLeuLeuGluLysAspGlyMetAlaPheGlnGlu
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Skeiky, Yasir A. W.
Fanger, Gary R.
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Smith, Carole L.
King, Gordon E.
Wang, Aijun
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Best Local Similarity:
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                                                             324 GCCCTAGACCCAGGGCCCTTTGACCAGGCAGCCCCTTTGCCCAGGAGCTGCTGGACGAC
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384 TCTGATGAGCTCAGTTGGATCATTGAGCTGCTGGAGAAGGATGGCATGGCCTTCCAGGAG 325
                                    153 AlaLeuAspProGlyProPheAspGlnGlySerProPheAlaGlnGluLeuLeuAspAsp 172
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
APPLICANT: Stoilk, John A.
APPLICANT: Stoilk, John A.
APPLICANT: Meagher, Madeleine J.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: CONFOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210121.411212
CURRENT PLLING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Mindows Version 3.0
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                                                                                                                  173 GlyGlnGlnAlaSerProTyrHisProGlySerCysGlyAlaGlyAlaProSerProGly
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US-09-833-263-853
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233 ArgaspCysLysGlyaspProLysHisGlyLysArgLysArgGlyArgProArgLys 252
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384 reneardadecreaerregarearreadecreergaagaagaagaregearregeerreergaag 325
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                                                        153 AlaLeuaspProGlyProPheAspGlnGlySerProPheAlaGlnGluLeuLeuAspAsp
                                                                                                                173 GlydlnGlnAlaSerProTyrHisProGlySerCysGlyAlaGlyAlaProSerProGly
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210.121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 853
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Xu, Jiangchun APPLICANT: Lodes, Michael J. APPLICANT: Secrist, Heather APPLICANT: Benson, Darin R. APPLICANT: Maspher, Madeleine Joy APPLICANT: Stolk, John A. APPLICANT: Wang, Tongtong
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624 GGTACAGAGAAGGCCAGCTGGTTGGGGGAACAGCCCCAGTTCTGGTCGAAGACGCGAGGTT 565
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       324 GCCCTAGACCCAGGCCCCTTTGACCAGGCAGCCCCTTTGCCCAGGAGCTGCTGGACGAC 265
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53 GlyThrGluLysAlaSerTrpLeuGlyGluGlnProGlnPheTrpSerLysThrGlnVal
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APPLICANT: Xu, Janaghun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Branch, John A.
APPLICANT: Mang, Yuqiu
APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Carole Lynn
APPLICANT: Ming, Addon E.
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Patent No. US20020076414A1
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ORGANISM: Homo sapiens
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Sequence 944, Application US/09833263

Sequence 944, Application US/09833263

Patent No. US20020110547A1

GENERAL INFORMATION:

APPLICANT: Wang, Aljun
APPLICANT: Clapper, Jonathan D.

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE

TITLE OF INVENTION: 10901-04-10

NUMBER OF SEQ ID NOS: 1093

SOUTHARE: FastESQ for Windows Version 3.0

SEQ ID NO 944

LENGTH: 563

TYPE: DNA

TYPE: DNA

ORGANISM: Homo sapien

US-09-833-263-944
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Perfect score:
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Description	Sequence 74,	Sequence 101	Sequence 563	Sequence 342	Sequence 192	Sequence 1105	Sequence 1105	Sequence 207,	Sequence 756	Sequence 64,	Sequence 853,	Seguence 853	Sequence 853,	Sequence 944,	Sequence 944.
ID	US-10-097-340-74	US-09-964-824A-101	US-09-964-824A-563	US-09-880-107-3420	US-09-967-768A-192	US-09-922-217-1105	US-10-025-380-1105	US-09-925-301-207	US-10-264-049-756	US-10-131-410-64	US-09-922-217-853	US-09-833-263-853	US-10-025-380-853	US-09-922-217-944	US-09-833-263-944
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US-10-025-380-944 US-09-664-287A-282 US-09-561-621-282 US-09-551-621-282 US-09-551-621-282 US-10-10-107-882-282 US-10-10-07-882-282 US-10-07-882-282 US-09-998-598-223 US-09-998-598-223 US-10-998-598-223 US-10-222-798-126 US-09-998-598-134 US-09-759-143-44 US-09-759-144-44 US-09-759-144-44 US-09-895-793-44 US-09-895-793-44 US-09-895-794-44 US-09-895-794-44 US-09-895-894-44 US-09-895-894-44 US-09-895-793-44 US-09-895-894-44 US-09-895-894-44 US-09-895-894-44 US-09-995-793-44 US-09-995-793-44 US-09-995-793-44 US-09-995-793-44 US-09-995-793-44 US-09-995-793-44 US-09-995-793-44 US-09-995-793-44 US-10-144-6784-44 US-10-10-940-44
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ALIGNMENTS

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TITLE OF INVENTION: Masessment, Prevention, and Therapy of Ovarian Cancer ITILE OF INVENTION: Masessment, Prevention, and Therapy of Ovarian Cancer TITLE OF INVENTION: Masessment, Prevention, and Therapy of Ovarian Cancer FILE REPERENCE: MRI-030

CURRENT APPLICATION NUMBER: US/10/097,340

CURRENT FILING DATE: 2002-03-14

PRIOR APPLICATION NUMBER: 60/276,025

PRIOR FILING DATE: 2001-03-14

PRIOR FILING DATE: 2001-03-14

PRIOR FILING DATE: 2001-03-14

PRIOR PILING DATE: 2001-03-14

PRIOR PILING DATE: 2001-03-14

PRIOR PILING DATE: 2001-09-26

PRIOR APPLICATION NUMBER: 60/311,732

PRIOR PILING DATE: 2001-09-26

PRIOR PILING DATE: 2001-09-26

PRIOR PILING DATE: 2001-09-26

PRIOR FILING DATE: 2001-09-26
              Sequence 74, Application US/10097340
Publication No. US2003008725041
GENERAL INFORMATION:
APPLICANT: John WONNHAN
APPLICANT: Sebastian HOERSCH
APPLICANT: Shubhangi KAWATKAR
APPLICANT: Steve G. KOVATS
APPLICANT: Rachel E. MEYERS
APPLICANT: Michael MORRISEY
APPLICANT: Michael MORRISEY
APPLICANT: Peter OLANDT
                                                                                                                                                                                                                                                                                                                                                                                   Gordon B. MILLS
Robert C. BAST, Jr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosemarie SCHMANDT
Xumei ZHAO
Karen GLATT
                                                                                                                                                                                                                                                                                                                                                         Peter VEIBY
                                                                                                                                                                                                                                                                                                                                                                                                                                    Karen LU
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US-10-097-340-74
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APPLICANT:
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APPLICANT:
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361 CICCATGCCCAGCTGCGAGACCTCACCTTCTGATGAGCTCAGTTGGATCATT 420
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Sequence 101, Application US/09964824A
; Patent No. US20020102531A1
; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 60/323,580
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 74
LENGTH: 1907
                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.
Matches 1116; Conservative
                                                                                                                                                                                                                   ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-097-340-74
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TILE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat TITLE OF INVENTION: Sets STILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT PILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
SOFTWARE PATENTING DATE: 2000-09-28
                                                                                                                                                                                                                                                                                                                 1021 ATCCTGGAACGGGTGGATGGCCGGCGACTCGTCTACAAGTTTGGCAAAAACTCAAGCGGC 1080
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Matches 1116; Conservative 0; Mismatches 0; Indels 0;
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FILE REFERENCE: 669290-73
CURRENT APPLICATION NUMBER: US/69/236, 033
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100.0%; Score 1116; DB 10; Length 1915;
Best Local Similarity 100.0%; Pred. No. 2e-313;
Matches 1116; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                              APPLICANT: Horne, Darci T.

APPLICANT: Horne, Darci T.

APPLICANT: Scherf, Joseph G.
APPLICANT: Gene Logic, Inc.
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer FILE REFERENCE: 4491-5028-W0
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT PILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 3420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Genbank Accession No. US20020142981A1 U73843
US-09-880-107-3420
                                                                                                                   1081 TGGAAGGAGGAAGATTCTCCAGAGTCGGAACTGA 1116
                                                                                                                                              1200 TGGAAGGAGGAAGGTTCTCCCAGAGTCGGAACTGA 1235
                                                                                                                                                                                                                                                          Sequence 3420, Application US/09880107
Patent No. US20020142981A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
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Sequence 192, Application US/09967768A
Patent No. US20020150877A1
GENERAL INFORMATION:
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signat
TITLE OF INVENTION: Cancer Sets
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Sequence 1105, Application US/09922217
; Patent No. US20020076414A1
; GENERAL INFORMATION:
                                                                                                                                                                                                     APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Becriet, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
APPLICANT: Jiang, Yuqiu
APPLICANT: Jiang, Yuqiu
                                                                                                                                                                                                                                                                                                                                          Jiang, Yuqiu
Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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ORGANISM: Homo sapiens
                                                                                                            RESULT 6
US-09-922-217-1105
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                                      1 ATGGCTGCAACCTGTGAGATTAGCAACATTTTTAGCAACTACTTCAGTGCGATGTACAGC 60
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                                                          120 ATGGCTGCAACCTGTGAGATAGCAACATTTTTAGCAACTACTTCAGTGCGATGTACAGC
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100.0%; Score 1116; DB 9; Length 1917;
Best Local Similarity 100.0%; Pred. No. 2e-313;
Matches 1116; Conservative 0; Mismatches 0; Indels 0;
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION: 08 COLON CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION NUMBER: US/09/922,217
CURRENT APPLIANCE DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1105
LENGTH: 1917
                              1200 TGGAAGGAGGAAGATTCTCCAGAGTCGGAACTGA 1235
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                                                                                                                                                 Query Match 100.0%; Score 1116; DB 14; Length 1917; Best Local Similarity 100.0%; Pred. No. 2e-313; Matches 1116; Conservative 0; Mismatches 0; Indels 0;
                          TYPE: DNA
CORGANISM: Homo sapiens
US-10-025-380-1105
   LENGTH: 1917
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APPLICANT: Lodes, Michael J.
APPLICANT: Bersist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
APPLICANT: Wing, Yugiu
APPLICANT: Smith, Carole L.
APPLICANT: King, Gordon E.
APPLICANT: King, Gordon E.
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APPLICANT: Smith, Carole L.
APPLICANT: Smith, Carole L.
APPLICANT: King, Gordon E.
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Panger, Gary R.
APPLICANT: Panger, Tomas S.
APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOUNDS FO
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Skeiky, Yasir A. W.
Fanger, Gary R.
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; Bedience 756, Application US/10264049
; Publication No. US20040005579A1
; GENERAL INFORMATION:
    APPLICANT BITSE et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE PA133P1
; CURRENT APPLICATION NUMBER: US/10/264,049
; PRIOR PELING DATE: 2002-10-04
; PRIOR PLLING DATE: 2001-06-07
; PRIOR PILING DATE: 2001-06-07
; PRIOR PILING DATE: 2000-06-07
; NUMBER OF SEQ ID NOS: 4360
; SOFTWARE: Patentin Ver: 3.1
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Matches 1112; Conservative
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Pred. No. 2.7e-313;
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FILE REFERENCE: PAIO (2007)
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PAICHIN VET: 2.0
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; Patent No. US20020052308A1
; GENERAL INFORMATION:
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APPLICANT: SPECHT, THOMAS
APPLICANT: HINAMAN, BERND
APPLICANT: HINAMAN, BERND
APPLICANT: SCHMITT, ARMIN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: DAHL, BDGAR
APPLICANT: DAHL, ANDRE
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMORS
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM TISSUE OF BREAST
TITLE OF INVENTION: TUMORS
TILLE REFERENCE: SCH-163
CURRENT APPLICATION NUMBER: US/10/131,410
FILE REPLICATION NUMBER: PCT/0E99/00908
PRIOR FILING DATE: 2000-09-20
PRIOR FILING DATE: 1999-03-19
NUMBER OF SEQ ID NOS: 202
SEQ ID NO 64
LENGTH: 2269
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                        US-10-131-410-64
; Sequence 64, Application US/10131410
; Publication No. US20030235915A1
; Publication INFORMATION:
; APPLICANT: SPECHT, THOMAS
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TCGAGGACTCCACCCTGGCCTCTGTTCCCCCTGCTGCCACCTTTGGGGCCCGATGACTTG 120
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                                      221 TCGGAGGACTCCACCCTGGCTCTGTTCCCCCTGCTGCCCCTTTGGGGGCCGATGACTTG
                                                                                                                                                                                      281 GTACTGACCCTGAGCAACCCCCAGATGTCATTGGAGGGTACAGAGAGGCCAGCTGGTTG
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335 TGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCCCAGCTGCGAGACCTCACTTCCAGCT 394
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; Patent No. US20020110547A1
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Best Local Similarity 99.88
Matches 625; Conservative
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CRGANISM: Homo sapien
US-09-833-263-853
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LENGTH: 626
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                                                                                                                                                                                                                                   TGGCCCAACTATGGGGCCCAAAAGAAAAAAAACAACAACATGACCTACGAGAAAGCTGAGCC 841
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                                                        812 GAGGCACCCACCTGTGGGAGTTCATCCGGGACATCCTCATCCACCCGGAGCTCAACGAGG 871
                                                                                             662 GAGGCACCCACCTGTGGGAGTTCATCCGGGACATCTTCATCCACCCGGAGCTCAACGAGG 721
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 853
LENGTH: 626
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Secrist, Heather
Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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Best Local Similarity 99.84
Matches 625; Conservative
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APPLICANT: Xu, Jiangchun
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ORGANISM: Homo sapiens
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APPLICANT:
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APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
APPLICANT: Capper, Jonathan D.
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REPERENCE: 210121.471012
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT PILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
NUMBER OF SEQ ID NOS: 1093
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99.8%; Pred. No. 6.5e-171;
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50.3%; Score 561.4;
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; ORGANISM: Homo sapiens
US-09-922-217-944
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210.121, 471.014
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
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55.9%; Score 624.4; DB 14;
Best Local Similarity 99.8%; Pred. No. 6.5e-171;
Matches 625; Conservative 0; Mismatches 1;
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SUFFRARE: FastSEQ for Windows Version 4.0
EQ OF NO 853
LENGTH: 626
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APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Onh A.
APPLICANT: Magg, Tongtong
APPLICANT: Wang, Tongtong
APPLICANT: Smith, Carole L.
APPLICANT: Smith, Carole L.
APPLICANT: King, Gordon E.
APPLICANT: King, Gordon E.
APPLICANT: King, Applicant Skeiky, Yasir A.
APPLICANT: Steiky, Yasir A.
APPLICANT: Steiky, Yasir A.
APPLICANT: Steiky, Yasir A.
APPLICANT: Vedyick Thomas S.
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ORGANISM: Homo sapiens
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                                                                                                         275 TCTCACGATGTGACATGGATGGCGCCACCCTCTGCAATTGTGCCCTTGAGGAGCTGCGTC
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
PILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 9; Length 563;
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 944
LENGTH: 663
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Patent No. US20020076414A1
GENERAL INFORMATION:
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Secrist, Heather
Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
Jiang, Yuqiu
Smith, Carole Lynn
King, Gordon E.
Wang, Aljun
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
APPLICANT: State Clapper, Jonathan D.
APPLICANT: State Clapper, Jonathan D.
APPLICANT: Madelaine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REPERBYCE: 210121.471C12
CURRENT APPLICATION NUMBER: US/09/833,263
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                                                                                                                                            219 GGACTGGATCAGCTACCAAGTGGAGAACAAGTACGACGCAAGCGCCATTGACTTCTC
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                                                                 Indels
     99.8%; Pred. No. 1.2e-152; vative 0; Mismatches 1;
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; Patent No. US20020110547A1
; GENERAL INFORMATION:
                                                  Matches 562; Conservative
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US-09-833-263-944
Best Local Similarity
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